

STIC Search Report Biotech-Chem Library

STIC Database Tracking Number: 194203

TO: Sheela Huff

Location: rem/3A15/3C18

Art Unit: 1643

Friday, June 30, 2006

Case Serial Number: 10/671054

From: Willis McCumber

Location: Biotech-Chem Library

REM-1A75

Phone: (571)272-2697

willis.mccumber@uspto.gov

Search Notes

Examiner Huff,

See attached results.

If you have any questions about this search feel free to contact me at any time.

Thank you for using STIC search services.

Willis McCumber ASRC Aerospace Corporation Library Technician STIC Biotech/Chem Library (571)272-2697



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STIC-Biotech/ChemLib

194203

From:

Huff, Sheela

Sent:

Tuesday, June 27, 2006 5:27 PM

To: Subject: STIC-Biotech/ChemLib 10671054--search request

Please search and interference search SEQ ID No. 1-2 of the above.

Thanks-

Sheela Huff
Art Unit 1643
571-272-0834
Remsen 3A15
mailbox Remsen 3C18

FOR

SCANNING ONLY

Searcher: ______ Searcher Phone: _____ Date Searcher Picked up: _____ Date completed: _____ Searcher Prep Time: _____ Online Time: _____ Online Time: _____

Type of Search

NA#______ AA#:_____

S/L:____ Oligomer:_____

Encode/Transl:_____ Text:____

Inventor:_____ Litigation:____

Vendors and cost where applicable STN:

DIALOG:
QUESTEL/ORBIT:
LEXIS/NEXIS:
SEQUENCE SYSTEM:
WWW/Internet:
Other (Specify):

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STIC SEARCH RESULTS FEEDBACK FORM

Biotech-Chem Library

Questions about the scope or the results of the search? Contact the searcher or contact:

Mary Hale, Information Branch Supervisor 571-272-2507 Remsen E01 D86

| /ol | untary Results Feedback Form |
|-----|--|
| > | I am an examiner in Workgroup: Example: 1610 |
| > | Relevant prior art found, search results used as follows: |
| | ☐ 102 rejection |
| | ☐ 103 rejection |
| | ☐ Cited as being of interest. |
| | ☐ Helped examiner better understand the invention. |
| | Helped examiner better understand the state of the art in their technology. |
| | Types of relevant prior art found: |
| | ☐ Foreign Patent(s) |
| | Non-Patent Literature (journal articles, conference proceedings, new product announcements etc.) |
| > | Relevant prior art not found: |
| | Results verified the lack of relevant prior art (helped determine patentability). |
| | Results were not useful in determining patentability or understanding the invention. |
| Co | amments: |

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A;Accession: A48089
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-673 «MAN»
A;Cross-references: UNIPROT:Q61592; UNIPARC:UPI000004BA5B; GB:X59846; NID:g407060; PIDN
A;Cross-references: UNIPROT:Q61592; UNIPARC:UPI000004BA5B; GB:X59846; NID:g407060; PIDN
A;Note: authors translated the codon CCC for residue 424 as Ile
C;Genetics:
A;Gene: gas6
C;Superfamily: plasma protein S; EGF homology; Gla domain homology; laminin G repeat how
F;38-89/Domain: Gla domain homology *status atypical *GLA*
F;117-150/Domain: EGF homology *EG1*
F;157-192/Domain: EGF homology *EG2*
F;198-233/Domain: EGF homology *EG3*
F;308-666/Domain: EGF homology *EG3*
F;308-666/Domain: EGF homology *EG3*
F;318-470/Domain: laminin G repeat homology *CGR*
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A48089
growth arrest-specific protein gas6 - mouse
C;Species: Mus musculus (house mouse)
C;Date: 26-May-1994 #sequence_revision 26-May-1994 #text_change
C;Accession: A48089; S37437
R;Manfioletti, G.; Brancolini, C.; Avanzi, G.; Schneider, C.
Mol. Cell. Biol. 13, 4476-44985, 1993
A;Title: The protein encoded by a growth arrest-specific gene (
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124
                     127 RKGTQACQDLMGNFFCLCKAGWGGRLCDKDVNECSQENGGCLQICHNKPGSFHCSCHSGF
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                                                                                                                                                                                                            PGPAALRRAPQLLLLLLAAECALAALLPAREATGFLRPRQRRAFQVFEEAKQGHLERECV
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:: Gla domain homology #Status atypical <GLA>
in: EGF homology <EGI>
:in: laminin G repeat homology <LGR>
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81.6%; Pred. No. 1.66
tive 45; Mismatches
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A;Molecule type: mRNA
A;Residues: 1-10, 'P', I25, 'L', 27-676 <HOS>
A;Crose-references: UNIPARC:UPI000014279E; GB:M15036; NID:g190288; PIDN:AAA36479.
R;Lundwall, A.; Dackowski, W.; Cohen, E.; Shaffer, M.; Mahr, A.; Dahlback, B.; St
Proc. Natl. Acad. Sci. U.S.A. 83, 6716-6720, 1986
A;Title: Isolation and sequence of the cDNA for human protein S, a regulator of b
                                                                                                                                     A;Cross-references: UNIPARC:UPI0000173343; GB:J02918 R;HOSKINS, J.; Norman, D.K.; Beckmann, R.J.; Long, G.L. Proc. Natl. Acad. Sci. U.S.A. 84, 349-353, 1987 A;Title: Cloning and characterization of human liver cDNA A;Reference number: A26157; MUID:87092407; PMID:3467362 A;Accession: A26157
                                                                                                                                                                                                                                                                                                                         A;Molecule type: DNA
A;Residues: 1-676 <SCH>
A;Residues: 1-676 <SCH>
A;Residues: 1-676 <SCH>
A;Residues: GB:M57853; NID:g190547;
A;Cross-references: UNIPROT:P07225; UNIPARC:UPI00001323E8; GB:M57853; NID:g190547;
A;Note: the authors translated the codon TTT for residue 26 as Leu
R;Ploos van Amstel, H.K.; Reitsma, P.H.; van der Logt, C.P.E.; Bertina, R.M.
Biochemistry 29, 7853-7861, 1990
A;Title: Intron-exon organization of the active human protein S gene PSalpha and it
A;Reference number: A35611; MUID:91084445; PMID:2148111
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N;Alternate
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                                                                                                                                                                                                                                                                                A; Molecule type: DNA
A; Residues: 1-25 < PL3>
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;Alternate names: vitamin K-dependent protein
;Species: Homo sapiens (man)
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F;25-41/Domain: propertide #status predicted <PRO>
F;26-85/Domain: Gla domain homology <GLA>
F;42-85/Domain: Gla domain homology <GGLA>
F;42-85/Domain: EGF homology <EG1>
F;111-154/Domain: EGF homology <EG2>
F;111-154/Domain: EGF homology <EG3>
F;161-199/Domain: EGF homology <EG3>
F;247-282/Domain: EGF homology <EG3>
F;247-282/Domain: EGF homology <EG3>
F;247-282/Domain: EGF homology <EG4>
F;315-667/Domain: EGF homology <EG4>
F;315-667/Domain: EGF homology <EG4>
F;315-67/Domain: EGF homology <EG4
F;317,60,61,61,67,70,73,77/Modified site: Expension of Egg4
F;317,11-112/Cleavage site: Agf-3145-154,161-175,171-184,186-199,205-217,212-226,228-241,
F;315-Modified site: Expthro-beta-hydroxyaspartic acid (Asp) #status predicted
F;317,219,258/Modified site: expthro-beta-hydroxyaspartic acid (Asp) #status predicted
F;319,530/Binding site: carbohydrate (Asp) (covalent) #status predicted
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C; Complex: :
C; Function:
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A;Accession: A25891
A;Molecule type: mRNA
A;Residues: 27-220,'L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Description: a cofactor for activated protein C (EC 3.4.21.69); thrombin cleavage des C;Superfamily: plasma protein S; EGP homology; Gla domain homology; laminin G repeat how C;Keywords: beta-hydroxyaspartic acid; blood coagulation; carbo F;1-24/Domain: signal sequence #status predicted <SIG>
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R;Ploos van Amstel, H.K.; van der Zanden, A.I
FEBS Lett. 222, 186-190, 1987
A;Title: Human protein S cDNA encodes Phe-16
A;Reference number: S02424; MUID:88005138; PM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Cross-references: GDB:120721; OMIM:176880
A;Map position: 3p11.1-3q11.2
A;Introns: 26/1; 78/3; 87/1; 116/1; 157/1; 201/1; 243/1; 283/3; 322/2; C;Complex: in plasma forms a complex with C4b binding protein
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R;Ploos van Amstel, J.K.; van der Zanden, A.L.; Bakker,
Thromb. Haemost. 58, 982-987, 1987
A;Title: Two genes homologous with human protein S cDNA
A;Reference number: A60903; MUID:88178564; PMID:2895503
A;Accession: A60903
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A;Title: Molecular analysis of the gene for vitamin K dependent pro A;Title: Molecular analysis of the gene for Vitamin K dependent pro A;Reference number: A35612; MUID:91084446; PMID:2148112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Gene: GDB: PROS1; PROS
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A; Residues: 1-676 < PL2 >
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A; Residues: 284-676 <EDE>
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A;Residues: 351-676 <PLO>
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Best Local S
Matches 298
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                                                                                                                                                                                                                                                                                                                                 18 LILLILLAAECALAALLPAREATQFLRPRQRRAFQVFEEAKQGHLERECVEELCSREEARE
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                        VFENDPETDYFYPRYLDCINKY-----GSPYTKNS--GFATCVQNLPDQCTPNPCDRKG
YMSCKDGKASFTCTCKPGWQGEKCEFDINECKDPSNINGGCSQICDNTPGSYHCSCKNGF
                                                             TQACQDLMGNFFCLCKAGWGGRLCDKDVNEC---SQENGGCLQICHNKPGSFHCSCHSGF
                                                                                                                                  VFENDPETDYFYPKYLVCLRSFQTGLFTAARQSTNAYPDLRSCVNAIPDQCSPLPCNEDG
                                                                                                                                                                                                                                                                           LACLLLVLPVSEANFLSKQQASQVL-VRKRRANSLLEETKQGNLEREC1EELCNKEEARE
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44.4%;
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Pred. No. 2e-100;
7; Mismatches 21
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PMID:2820795
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М.
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A;Cross-references: UNIPARC:UPI0000176513; EMBL:L31380
A;Experimental source: tissue type liver
A;Note: the source is designated as rhesus monkey
C;Genetics:
A;Gene: PROS
C;Superfamily: plasma protein S; EGF homology; Gla doma
F;1-51/Domain: Gla domain homology (frame)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       plasma protein S precursor, vitamin K dependent - rhesus macaque (fragment) (Species: Macaca mulatta (rhesus macaque) (plate: 19-Mar-1997 #sequence_revision 18-Jul-1997 #text_change 16-Jul-1999 (Accession: $53434 R;Greengard, J.S.; Fernandez, J.A.; Radtke, K.P.; Griffin, J.H. Biochem. J. 305, 397-403, 1995 A;Title: Identification of candidate residues for interaction of protein S 1A;Reference number: $53433; MUID:95134217; PMID:7832752
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                                                                                                                                                                                             C;Superfamily: plasma protein S; EGF homology; Gla domain homology; laminin F;1-51/Domain: Gla domain homology (fragment) <GLA>F;1-7/Domain: signal sequence (fragment) #status predicted <SIG>F;8-642/Product: plasma protein S #status predicted <MAT>F;8-120/Domain: EGF homology <EG1>F;87-120/Domain: EGF homology <EG2>
                                                                                                                                                                                                                                                                                                                                                                                                                   A; Molecule type: mRNA
A; Residues: 1-642 < GRE>
                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Status: nucleic acid sequence not shown; not
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                                       Query Match
Best Local Similarity
Matches 287; Conserv
                                                                                                                                                          213-248/Domain:
                                                                                                                                                                               171-207/Domain:
                                                                                                                       291-444/Domain:
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  RQRRAFQVFEEAKQGHLERECVEELCSREEAREVFENDPETDYFYPRYLDCINKYGSPYT
                                     Conservative 122; Mismatches
                                                                                                                                   EGF homology <EG3>
EGF homology <EG4>
sex hormone-binding globulin homology
                                                                                                                   laminin G repeat homology <LGR>
                                                         41.2%;
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                                                         Score 1512.5; DB 2;
Pred. No. 2.6e-98;
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                                         Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RLCDKDVNEC---SQENGGCLQICHNKPGSFHCSCHSGFELSSDGRTCQDIDECA-DSEA
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                                                                                      LMEIKVCDGQEHVVTVSLRDGEATLEVDGTRGQSEVSAAQLQERLAVLERHLRSPVLTFA
                                                                                                                  WHINVTLNIRPSTGTGVMLALVSSN-NTVPFAVSLVD--STSE-KSQDIVISVENTVIYR
                                                                                                                                           WEVEYVAHIRPAADTGVLFALWAPDLRAVPLSVALVDYHSTKKLKKQLVVLAVEHTAL-A
                                                                                                                                                                                                      RSWNWLNGEDTTIQETVKVNTRMQCFSVTERGSFYPGSGFAFYSLDYMRTPLDVGTEST-
                                                                                                                                                                                                                                                                                               TFDPEGILLFAGGHODSTWIVLALRAGRLELQLRYNGVGRVTSSGPVINHGMWQTISVEE
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                                                                                                                                                                                                                                                                                                                                                                                                                                   CGEARCKNILPGSYSCLCDEGFAYSSQEKACRDVDECLQGRCEQVCVNSPGSYTCHCDGRG
GGLPDVPFSATPVNAFYNGCMEVNINGVELDLDEAISKHNDIRAHSCPSV
                                                         IQALSLCSYORS --HLEFRVNRNNLELLTPLKIETISQEELQTQLAILDKAMKGKVATYL
                                                                                                                                                                            RSWNLMKQGASGIKEIIQEKQNKHCLVTVEKGSYYPGSGIAEFHIDY-----NNGSNAEG
                                                                                                                                                                                                                                       LEHSISIKIAKEAVMDINKPGPLFKPENGLLETKVYFAGFPRKVESELIKPINPRLDGCI
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RESULT 6

KXBOS

plasma protein S precursor - bovine
N,Alternate names: vitamin K-dependent protein S
C;Species: Bos primigenius taurus (cattle)
C;Date: 08-Aug-1987 #sequence_revision 26-Jan-1996 #text_change
C;Accession: A24759, A23888
R;Jahlback, B; Lundwall, A.; Stenflo, J.
Proc. Natl. Acad. Sci. U.S.A. 83, 4199-4203, 1986
A;Title: Primary structure of bovine vitamin K-dependent protein
A;Reference number: A24759; MUID:86233400; PMID:2940598
A;Rocession: A24759
A;Rocession: A24759
A;Rocession: A24759
A;Rocession: A24759
A;Rocession: A24759
A;Rote: parts of this sequence, including the amino end of the
A;Roticlues: 1-675 < DAHY
A;Roticlues: 1-675 < DAHY
A;Roticlues: 1-675 < DAHY
A;Roticlues: 1-675 < DAHY
A;Rote: parts of this sequence, including the amino end of the
A;Roticlues: 1-675 < DAHY
A;Roticlues: 1-675 
                       on: a cofactor for activated protein C (EC 3.4.21.69); thrombin cleavage desuly: plasma protein S; EGF homology; Gla domain homology; laminin G repeat how beta-hydroxyasparagine; beta-hydroxyaspartic acid; blood coagulation; carbo-
                                                                                                                                                                                                                                                         C4b binding protein
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#text_change 09-Jul-2004

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,315-666/Domain: sex hormone-binding globulin homology <SHB>
,315-666/Domain: laminin G repeat homology <LGR>
,325-478/Domain: laminin G repeat homology <LGR>
,48,55,57,60,61,66,67,70,73,77/Modified site: gamma-carboxyglutamic acid (Glu) #stat
,47,48,55,57,60,61,66,67,70,73,77/Modified site: gamma-carboxyglutamic acid (Glu) #stat
,58-63,88-113,121-134,126-143,145-154,161-175,171-184,186-199,205-217,212-226,228-241,2
,93-94/Cleavage site: Arg-Ala (thrombin) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-41/Domain: propeptide #status predicted <PRO>
26-85/Domain: Gla domain homology <GLA>
42-675/Product: plasma protein S #status experimental
121-154/Domain: EGF homology <EG1>
161-199/Domain: EGF homology <EG2>
1205-241/Domain: EGF homology <EG3>
1247-282/Domain: EGF homology <EG4>
1315-666/Domain: EGF homology <EG4>
1315-666/Domain: sex hormone-binding globulin homology
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                                                                                                                                                                                                    YHSTKKLKKQLVVLAVEHTALALME-IKVCDGQEHVVTVSLRDGEATLEVDGTRGQSEVS
                                                                                                                                                                                                                                                                          GSGFAFYSLDYMRTPLDVGTESTWEVEVVAHIRPAADTGVLFALWAPDLRAVPLSVALVD
                                                                              SEDFOROFAILDEAMKGTVVTYLGGLPDVPFSATPVNAFYQGCMEVNINGVQVDLDEAIS
                                                                                                                                                                                                                                                                                                                                FAGVPRKMENALIRPINPRLDGCIRGWNLMNQGTSGVKEIIQEKQNKHCLVNVEKGSYYP
                                                                                                                                                                                                                                                                                                                                                                     VGGIPFH-EKDLVQPINPRLDGCMRSWNWLNGEDTTIQETVKVNTRMQCFSVTERGSFYP
                                                                                                                                                                                                                                                                                                                                                                                                                   KTTKMTTGGKVINDGLWHMVSVEELEQSISVKIAKEAVMNINKPGSLFKPTNGFLETKVY
                                                                                                                                                                                                                                                                                                                                                                                                                                                      GVGRVTSSGPVINHGMWQTISVEELARNLVIKVNRDAVMKIAVAGDLFQPERGLYHLNLT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GV-VLYLKFRLPETTRFSAEFDFRTYDSEGVILYAESSDHSAWFLIALREGKIEIQFKNE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AENICAQICVNYPGGYSCYCDGKKGFKLAQDQKGCEAVPVCLPLDLDKNYELLYLAEQFV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SGFELSSDGRTCQDIDECA-DSEACGEARCKNLPGSYSCLCDEGFAYSSQEKACRDVDEC 242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AREVFENDDETDYFYPRYLDCINKYGSPYTKNSGFAT------CVQNLPDQCTPNPCD 126
KHNDIRAHSCPSV 668
                                       KHSDITAHSCPPV 673
                                                                                                                   AAQLQERLAVLERHLRSPVLTEAGGLPDVPVTSAPVTAFYRGCMTLEVNRRLLDLDEAAY
                                                                                                                                                                                                                                                  GTGVAQFSINYK----NESNPEAWQINVSLNIRPSAGTGVMLALVSDN--TVPFALSLVD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GTPVIRLRFKRLQPTRLVAEFDFRTFDPEGILLFAGGHQDSTWIVLALRAGRLELQLRYN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NGFVMLSNKKDCKDVDECVLKPSICGTÁVCKNIPGDFECECAEGYKYNPVSKSCDDVDEC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EDGFWTCKDGQATFTCICKSGWQGEKCESDINECKDPVNINGGCSQICENTPGSYHCSCK 187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RKGTQACQDLMGNFFCLCKAGWGGRLCDKDVNECSQE---NGGCLQICHNKPGSFHCSCH 183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LLAAECALAALLPAREATQFLRP------RQRRAFQVFEEAKQGHLERECVEELCSREE
                                                                                                                                                               -SATEKL--QDILVSVESMVIGRIEAISLCSDQQ--TFLEIRVNRNNLELSTQLRKDSFH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LQGRCEQVCVNSPGSYTCHCDGRGGLKLSQDMDTCEDILPCVPFSVAKSVKSLYLGRMFS
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C;Superfamily:
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Best Local
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plasma protein S precursor, vitamin K dependent - process Sus scrofa domestica (domestic pig) C;Date: 01-Aug-1995 #sequence_revision 18-Jul-1997 C;Accession: S53433 R;Greengard, J.S.; Fernandez, J.A.; Radtke, K.P.; Chieffeld A.S.; Chieffeld A.S.; Radtke, K.P.; Chieffeld A.S.; Chieffeld
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F;87-120/Domain: EGF homology <EG1>
F;127-165/Domain: EGF homology <EG2>
F;171-207/Domain: EGF homology <EG3>
F;171-207/Domain: EGF homology <EG3>
F;213-248/Domain: EGF homology <EG4>
F;213-248/Domain: sex homology <EG4>
F;213-444/Domain: laminin G repeat homology <LGR>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Biochem. J. 305, 397-403, 1995
A;Title: Identification of candidate residues for interaction A;Reference number: S53433; MUID:95134217; PMID:7832752
A;Accession: S53433
A;Status: nucleic acid sequence not shown; not compared with c
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                                                                                                  DGQEHVVTVSLRDGEATLEVDGTRGQSEVSAAQLQERLAVLERHLRSPVLTFAGGLPDVP
SSQ--LSLLEMRVSVNSLELLTQFEKRRISSEDYQGQFAILDKAMEGTVATYLGGLPDVP
                                                                                                                                                                                                                                                                                             HIRPAADTGVLFALWAPDLRAVPLSVALVDYHSTKKLKKQLVVLAVEHTALA-LMEIKVC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IKVNRDAVMKIAVAGDLFQPERGLYHLNLTVGGIPFH-EKDLVQPINPRLDGCMRSWNWL
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                                                                                                                                                                                                                                                                                                                                                                                                                                              NQGASGVKEIIQEIQNKHCLVTVEKGSYYPGSGVAQFSIDYK----NVSSAGAWQVNVSL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NGEDTTIQETVKVNTRMQCFSVTERGSFYPGSGFAFYSLDYMRTPLDVGTESTWEVEVVA
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RESULT 8
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$C; Species: Oryct
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F;92-125/Domain: EGF homology <EGI>F;92-125/Domain: EGF homology <EGI
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C;Date: 18-Feb-1994 #sequence revision 21-Jul-1995 #text_change
C;Accession: S38819; S37238
R;He, X.; Dahlbaeck, B.
Eur. J. Biochem. 217, 857-865, 1993
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F;176-212/Domain: EGF homology <EG3>
F;218-253/Domain: EGF homology <EG4>
F;286-637/Domain: sex hormone-binding globulin homology
F;296-449/Domain: laminin G repeat homology <LGR>
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A; Residues: 1-502, 'L', 504-646 <HE2>
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A;Cross-references: UNIPROT:P98118; UNIPARC:UPI0000176514;
R;He, X.; Dahlbock, B.
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A;Title: Structure of mouse protein S as determined by PCR amplification and DNA sequenc A;Reference number: 159616; MUID:94302659; PMID:8029814
A;Recession: 159616
A;Reference number: 159616; MUID:94302659; PMID:8029814
A;Recession: 159616
A;Cccession: 159616
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Anolecule type: mRNA
A;Residues: 33-492,'L',494-675 <RES>
A;Cross-references: UNIPARC:UPI000016CFD2; GB:L27439; NID:9487866; PIDN:AAA40006.1; PID
C;Complex: in plasma forms a complex with C4b binding protein
C;Function:
A;Description: a cofactor for activated protein C (EC 3.4.21.69); thrombin cleavage dest
C;Superfamily: plasma protein S; EGF homology; Gla domain homology; laminin G repeat how
C;Superfamily: plasma protein S; EGF homology; Gla domain homology; laminin G repeat how
C;Keywords: beta-hydroxyasparagine; beta-hydroxyaspartic acid; blood coagulation; carbox
F;1-24/Domain: signal sequence #status predicted <SIG>
F;25-41/Domain: Gla domain homology <EG1>
F;26-85/Domain: Gla domain homology <EG1>
F;121-154/Domain: EGF homology <EG2>
F;121-154/Domain: EGF homology <EG3>
F;161-199/Domain: EGF homology <EG3>
F;161-199/Domain: EGF homology <EG3>
F;161-199/Domain: EGF homology <EG3>
F;161-199/Domain: EGF homology <EG4>
F;161-199/Domain:
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A;Title: Cloning and sequencing of a cDNA encoding the murine A;Reference number: $43504; MUID:94198297; PMID:8148380
A;Recession: $43504
A;Residnes: 1
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C;Date: 13-Jan-1995 #sequence_revision
C;Accession: S43504; I59616; S35962
R;Chu, M.D.; Sun, J.; Bird, P.
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F;47,48,55,57,60,61,66,67,70,73,77/Modified site: gamma-carboxyglutamic acid (Glu) #state;58-63,88-t13,121-134,126-143,145-154,161-175,171-184,186-199,205-217,212-226,228-241,711-112/Cleavage site: Arg-Ser (thrombin) #status predicted
F;131-112/Cleavage site: erythro-beta-hydroxyaspartic acid (Asp) #status predicted
F;177,219,258/Modified site: erythro-beta-hydroxyaspartic acid (Asp) #status predicted
F;177,219,258/Modified site: erythro-beta-hydroxyasparagine (Asn) #status predicted
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CDRKGTQACQDLMGNFFCLCKAGWGGRLCDKDVNEC---
                                                                                                                                            EEAREVFENDPETDYFYPRYLDCINKY--GSPYTKNSG-
                                                                                                                                                                                                                                               VLLA---CLALVIPVSE-TNPLSKERASQVLVRKRRANTLFEETMKGNLERECIEELCNK
                                                                                                          <u>ĖĖARĖVFĖNNPĖTDYFYPKYLGĆLGAFRVGSFHAARQSANAYPDLRSČVKAISDQČDPIP</u>
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43.0%;
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Pred. No. 4
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-SQENGGCLQICHNKPGSFHCS
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PLASMATS

PLASMA PROTEIN S precursor - rat

plasma protein S precursor - rat

Claspecies: Rattus norvegicus (Norway rat)

Claste: 04-Oct-1995 #Bequence_revision 26-Jan-1996 #text_change 09-Jul-2004

Claste: 04-Oct-1995 #Bequence_revision 26-Jan-1996 #text_change 09-Jul-2004

Claste: 04-Oct-1995 #Bequence_revision 26-Jan-1996 #text_change 09-Jul-2004

Claste: 04-Oct-1995 #Bequence, K.; Nishioka, J.; Suzuki, K.

R; Yasuda, F.; Hayashi, T.; Tanitame, K.; Nishioka, J.; Suzuki, K.

J. Blochem. 117, 374-383, 1995

A, Filtle: Molecular cloning and functional characterization of rat plasma protein S.

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A, Filtle: Molecular cloning and functional characterization of rat plasma protein S.

A, Filtle: Molecular cloning set and VIIIa fils protein that has an anticos activation of coagulation factors Va and VIIIa fils characterization of coagulation factors Va and VIIIa fils characterization of coagulation for set and part protein S.

A, Filtle: Molecular cloning set and VIIIa fils protein also binds to factor Va and Clauper family plasma protein S.

Filtle: Molecular cloning set and VIIIa fils characterization acid; blood coagulation; carbox Fils-Molecular protein S.

Filtle: Molecular cloning set and VIIIa fils cloning set and cloning set and clone set and clon
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7,42-675/Froduct: Glasma protein S #status predicted <MAT>
7,48-116/Domain: thrombin-sensitive #status predicted <THS>
7,88-116/Domain: EGF homology <EG1>
7,121-134/Domain: EGF homology <EG2>
7,121-134/Domain: EGF homology <EG2>
7,205-241/Domain: EGF homology <EG3>
7,205-241/Domain: EGF homology <EG3>
7,205-241/Domain: EGF homology <EG3>
7,315-666/Domain: Sex hormone-binding globulin homology <SHB>
7,325-478/Domain: laminin G repeat homology <LGR>
7,325-478/Domain: laminin G repeat homology <LGR>
7,47,48,55,57,60,61,66,67,70,73,77/Modified site: gamma-carboxyglutamic /F,58-63,88-113,121-134,126-143,145-154,161-175,171-184,186-199,205-217,2
F,58-63,88-113,121-134,126-143,145-154,161-175,171-184,186-199,205-217,2
F,111-112/Cleavage site: Arg-Ser (thrombin) #status predicted
7,136/Modified site: erythro-beta-hydroxyaspartic acid (Asp) #status predicted
7,137,219,258/Modified site: erythro-beta-hydroxyasparagine (Asn) #status
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                 AAYKHSDITAHSCPPV
                                                     VIYSKDLORQLAILDKTMKGTVATYLGGVPDISFSATPVNAFYSGCMEVNINGVQLDLDE
                                                                                                                                                                                                                                                LTVGGIPFH-EKDLVQPINPRLDGCMRSWNWLNGEDTTIQETVKVNTRMQCFSVTERGSF
 AISKHNDIRAHSCPSV
                                                                             EVSAAQLQERLAVLERHLRSPVLTFAGGLPDVPVTSAPVTAFYRGCMTLEVNRRLLDLDE
                                                                                                                                  VDYHSTKKLKKQLVVLAVEHTALALME-IKVCDGQEHVVTVSL-RDGEATLEVDGTRGQS
                                                                                                                                                              YPGSGIAQFSIDYN----NVTNAEGWQINVTLNIRPSTGTGVMLALVSGD--TVPFALSL
                                                                                                                                                                                        YPGSGFAFYSLDYMRTPLDVGTESTWEVEVVAHIRPAADTGVLFALWAPDLRAVPLSVAL
                                                                                                                                                                                                                    IYFAGLPRKVESALIKPINPRLDGCIRGWNLMKQGALGAKEIVEGKQNKHCFLTVEKGSY
                                                                                                                                                                                                                                                                                            FSGTPVIRLRFKRLQPTRLVAEFDFRTFDPEGILLFAGGHQDSTWIVLALRAGRLELQLR
                                                                                                                                                                                                                                                                                                                                                                                                     ECLQGRCEQVCVNSPGSYTCHCDGRGGLKLSQDMDTCEDILPCVPFSVAKSVKSLYLGRM
                                                                                                                                                                                                                                                                                                                                                                                                                                            CKIGFAMLTNKKDCKDVDECSLKPSVCGTAVCKNIPGDFECECPNGYRYDPSSKSCKDVD
                                                                                                                                                                                                                                                                                                                                                                                                                                                           CHSGFELSSDGRTCQDIDECA-DSEACGEARCKNLPGSYSCLCDEGFAYSSQEKACRDVD
                                                                                                          VDSGSG---TSQDILVFVENSVAAHLEAITLCSEQPSQLKCNINRNG---
                                                                                                                                                                                                                                                                                                                                  FAGV-VLYLKFRLPDITRFSAEFDFRTYDSEGIILYAESLDHSNWLLIALREGKIEVQFK
                                                                                                                                                                                                                                                                                                                                                                                       ECSENTCAQLCVNYPGGYSCYCDGKKGFKLAQDQRSCEGIPVCLSLDLDKNYELLYLAEQ
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42.8%; Pred. No. 9.3e-94;
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RESULT 11

IS9618

protein S - rat (fragment)
C;Species: Rattus norvegicus (Norway rat)
C;Species: Rattus norvegicus (Norway rat)
C;Date: 02-Aug-1996 #sequence_revision 02-Aug-1996 #text_change 09-Jul-2004
C;Accession: IS9618
R;Jamison, C.S.; McDowell, S.A.; Marlar, R.A.; Degen, S.J.F.
Thromb. Res. 78, 407-419, 1995
A;Title: Developmental expression of protein C and protein S in the rat.
A;Reference number: IS9618; MUID:95389408; PMID:7660357
A;Accession: IS9618
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-346 <RES>
A;Cross-references: UNIPROT:Q62628; UNIPARC:UPI00000E6E01; EMBL:U06230; NID:g497116;
C;Superfamily: plasma protein S; EGF homology; Gla domain homology; laminin G repeat
F;1-337/Domain: laminin G repeat homology (fragment) <LGR>
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bol IId

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A;Cross-references: UNIPROT:Q08879; UNIPARC:UPI0000176527; EMBL:X70854
R;Pan, T.C.; Kluge, M.; Zhang, R.Z.; Mayer, U.; Timpl, R.; Chu, M.L.
submitted to the EMBL Data Library, January 1993
A;Description: Sequence of extracellular mouse protein BM-90/fibulin and its calcium-dep
A;Reference number: S36440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R;Pan, T.C.; Kluge, M.; Zhang, R.Z.; Mayer, U.; Timpl, R.; Chu, Eur. J. Biochem. 215, 733-740, 1993
A;Title: Sequence of extracellular mouse protein BM-90/fibulin a A;Reference number: S34968; MUID:93358897; PMID:8354280
A;Recession: S78040
                                                                                                                                                                                                                                                                                   A;Molecule type: mRNA
A;Residues: 1-39, 'P',41-685 < CHU>
A;Cross references: UNIPARC:UPI000002A96F; EMBL:X70854
C;Genetics: A;Introns: 568/3
A;Introns: 568/3
C;Superfamily: fibulin-1; EGF homology
C;Keywords: alternative splicing; basement membrane; c:
C;Keywords: alternative splicing; basement decirce < CSIG>
F;10-29/Domain: signal sequence #status predicted < CSIG>
F;10-29/Domain: fibulin, splice form C #status predicted < CSIG>
F;98,537,541/Binding site: carbohydrate (Asn) (covalen)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 12
$78040
$78040
$100111, splice form C precursor - mouse
fibulin, splice form C precursor - mouse
fibulin, splice form C precursor - mouse
N;Alternate names: basement-membrane protein BM-90
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Species: Acul-1998 #sequence revision 24-Jul-1998 #text_change 09-Jul-2004
C;Accession: $78040; $78560; $36440
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A; Residues: 1-685 < PAN>
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                                                                                                                                                                                        Query Match
Best Local S
Matches 84
                                                                                                                                                                                                                                                                                   ;Reywords: alternative splicing; basement membrane; calcium binding;
;1-29/Domain: signal sequence #status predicted <SIG>
;30-685/Product: fibulin, splice form C #status predicted <MAT>
;98,537,541/Binding site: carbohydrate (Asn) (covalent) #status predi
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Best Local Similarity
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                                          135 DIMGNEFCLCKAGWG----GRICDKDVNECSQENGG-CLQICHNKPGSEHCSCHSGEBIS
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                                                                                                                                                                                                                 Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ALALME-IKVCDGQEHVVTVSL-RDGEATLEVDGTRGQSEVSAAQLQERLAVLERHLRSP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAEDWQINVTLNIRPFTGTGVMLALVSGD--TVPFALSLVDSGSG---TSQDILVFVENS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LDGCMRSWNWLNGEDTTIQETVKVNTRMQCFSVTERGSFYPGSGFAFYSLDYMRTPLDVG 500
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EFDFRTFDPEGILLFAGGHQDSTWIVLALRAGRLELQLRYNGVGRVTSSGPVINHGMWQT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VSVEELDDSVSIKIAKEAVMNINKLGSLFKPTDGFLDTKIYFAGLPRKVESALIKPINPR
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                                                                                                                                           CINKYGSPYTKNSGFATCVQNLP-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            VATYLGGVPDISFSATPVNAFYSGCMEVNINGVQLDLDEAISKHNDIRAHSCPSV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LDGCIRGWNLMKQGALGAKEIVEGKQNKHCFLTVEKGSYYPGSGIAQFSIDYN----NVT 172
NSPGSFRCECKAGFYFDGISRTC-VDINECQRYPGRLCGHKCENTPGSFHCSCSAGFRLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                           Conservative
                                                                                                                                                                                                               9.9%;
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                                                                                             ---YTCQKNVPNCGRGYHLNEEGTRCVDVDECAPPAEPCG-KG-HHCL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      71;
                                                                                                                                                                                        24; Mismatches
                                                                                                                                                                                                               Pred.
                                                                                                                                                                                                                 Score 362; DB 2;
Pred. No. 1.5e-17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          -LELWTPVRKDVIYSKDLQRQLAILDKTMKGT
                                                                                                                                                                                           68;
                                                                                                                                                                                                                                      Length 685
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RESULT A47221

C;Species:

Homo

sapiens

human (man)

(fragment)

fibrillin 1 precursor

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A; Molecule type: mRNA
A; Residues: 1-39, 'p', 41-705 < PAW>
A; Experimental source: cell-line F9 teratocarcinoma
R; Kluge, M.; Mann, K.; Dziadek, M.; Timpl, R.

Bur, J. Biochem. 193, 651-659, 1990
A; Title: Characterization of a novel calcium-binding 90-kDa glycoprotein (BM-90) shared
A; Reference number: S13814
A; Accession: S13814
A; Accession: S13814
A; MOLD: 91065369; PMID: 2249686
A; Residues: 28; 31-49, 'X', 51-53; 'XX', 110-117; 231-240, 'X', 242-243; 339-362, 'S', 364-387; 434-A; Cross-references: UNIPARC: UPI0000176520, 525; UNIPARC: UPI0000176520
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Cross-references: UNIPROT:Q08879; UNIPARC:UPI000017651E
R;Pan, T.C.; Kluge, M.; Zhang, R.Z.; Mayer, U.; Timpl, R.; Chu, M.L.
submitted to the EMBL Data Library, January 1993
A;Description: Sequence of extracellular mouse protein BM-90/fibulin
A;Reference number: S36440
A;Accession: S36441
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C;Species: Mus musculus (house mouse)
C;Date: 10-Dec-1993 #sequence revision 10-Nov-1995 #text_change 09-Jul-2004
C;Accession: S34968; S36441; S13814
R;Pan, T.C.; Kluge, M.; Zhang, R.Z.; Mayer, U.; Timpl, R.; Chu, M.L.
Bur. J. Biochem. 215, 733-740, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eur. J. Blochem. 215, 733-740, 1993
A;Title: Sequence of extracellular mouse protein BM-90/fibulin and its
A;Reference number: S34968; MUID:93358897; PMID:8354280
A;Accession: S34968
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Molecule type: mRNA
A; Residues: 1-705 < PAN>
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Superfamily: fibulin-1; EGF homology
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                                                                                                                                                                                                                                                                                                                                            327 CINTEGS-----YTCOKNVPNCGRGYHLNEEGTRCVDVDECAPPAEPCG-KG-HHCL
                                                                                                                                                                         190 SDGRTCQDIDECADSEACGEARCKNLPGSYSCLCDEGFAYSSQE-KACRDVDECLQGR--
                                                                                                                                                                                                                                  377 NSPGSFRCECKAGFYFDGISRTC-VDINECORYPGRLCGHKCENTPGSFHCSCSAGFRLS
                                                                                                                                                                                                                                                                                   135 DLMGNFFCLCKAGWG----GRLCDKDVNECSQENGG-CLQICHNKPGSFHCSCHSGFELS
                                                                                                                                                                                                                                                                                                                                                                                                95 CINKYGSPYTKNSGFATCVQNLP----
                                                                                                                                                                                                                                                                                                                                                                                                                                                         84;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HICSYRCINIPGSFQCSCPS-SGYRLAPNGRNCQDIDECV
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HICSYRCINIPGSFOCSCPS-SGYRLAPNGRNCQDIDECV
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                                                               --CEQVCVNSPGSYTCHCDGRGGLKLSQDMDTCEDILPCV
                                                                                                                        VDGRSCEDVNECLNSPCSQE--CANVYGSYQCYCRRGYQLSDVDGVTCEDIDECALPTGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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Pred. No. 1.5e-17;
P4; Mismatches 68
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A;Accession: S62111
A;McCession: S62111
A;McCession: S62111
A;McCession: S72111
A;McCession: S72111
A;McCession: S72111
A;McCession: S72111
A;McCession: S72111
A;McCession: S72111
Biol. Chem. 264, 21381-21385, 1989
A;Title: Connective tissue microfibrils. Isolation and ch A;Reference number: A34198; MUID:90078246; PMID:2512293
A;Accession: A34198
A;Accession: A34198
                                                                                                                                                                                                                              A;Molecule type: DNA
A;Residues: 2217-2288,'I',2290-2325 <RES>
A;Residues: 2217-2288,'I',2290-2325 <RES>
A;Cross-references: UNIPARC:UPI00006FPF6; GB:S54426; NID:g264860; PIDN:AAB25244.1; P.
A;Cross-references: UNIPARC:UPI00006FPF6; GB:S54426; NID:g264860; PIDN:AAB25244.1; P.
R;Lee, B.; Godfrey, M.; Vitale, E.; Hori, H.; Mattei, M.G.; Sarfarazi, M.; Tsipouras,
Nature 352, 330-334, 1991
A;Title: Linkage of Marfan syndrome and a phenotypically related disorder to two difference number: S17062; MUID:91304567; PMID:1852206
A;Accession: S17062
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hum. Mol. Genet. 2, ybi-ybo, xyzy
A;Title: Genomic organization of the sequence coding fo:
A;Reference number: I54355; MUID:93372860; PMID:8364578
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Accession: A47221; I54355; $\overline{S}17064; I59574; S17062; $62\overline{S}11; $\overline{A}34198} R;Corson, G.M.; Chalberg, S.C.; Dietz, H.C.; Charbonneau, N.L.; Sakai, L.Y. Genomics 17, 476-484, 1993
A;Title: Fibrillin binds calcium and is coded by cDNAs that reveal a multidomain A;Reference number: A47221; MUID:94010947; PMID:7691719
A;Accession: A47221
                                                                                                                                                                     F;132-3002/Product: f
F;1332-1367/Domain: E
F;1457-1492/Domain: E
F;2262-2295/Domain: E
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A; Residues: 1030-3002 <MAS>
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A;Title: Partial sequence of a candidate gene for the Marfan syndrome.
A;Reference number: S17064; MUID:91304568; PMID:1852207
A;Accession: S17064
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A;Molecule type: mRNA
A;Residues: 132-3002 <PER>
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A;Cross-references: UNIPROT:P35555; UNIPROT:Q15972;
R;Pereira, L.V.; D'Alessio, M.; Ramirez, F.; Lynch,
Hum. Mol. Genet. 2, 961-968, 1993
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        Query Match
Best Local S
Matches 88
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;Molecule type: mRNA
Residues: 'VLVTVVFIFLSYNKML',944-1444 <LEE1>
;Cross-references: UNIPARC:UPI0000072A79; EMBL:X62008; NID:g31398;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Status: preliminary; translated from GB/EMBL/DDBJ
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             ; 88
                                           Similarity
    9.6%;
larity 30.6%;
Conservative 2
                                                                                                                                                                                                        fibrillin (5'-region exon
EGF homology <EGF>
EGF homology <EGF2>
                                                                                                                                                                         EGF
                                                                                                                                                                 homology
        26;
                                                                                                                                                                     <EGF1>
Score 352; DB 2;
Pred. No. 4.5e-16;
6; Mismatches 78;
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                                                                          Length 3002;
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Indels
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96;
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Gaps
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Search completed: Job time : 46 secs

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A;Title: Sequence of the coding region of the bovine fibrillin A;Reference number: A55567; MUID:95137597; PMID:7835900
A;Accession: A55567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      fibrillin I - bovine
C;Species: Bos primigenius taurus (catt
C;Date: 20-Pe___1995 #sequence_revision
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Best Local (
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                             RCEQVCVNSPGSYTCHCD----GRGGLKLSQDMDTCE
                                                             SCTDIDECEDNPNICOGGOCTNIPGEYRCLCYDGFMASEDMKTCVDVNECDLNPNICLSG
                                                                                           TCQDIDECADS-EACGEARCKNLPGSYSCLCDEGFAYSSQEKACRDVDE-----CLQG
                                                                                                                        IGKYQCACNPGYHSTPDRLFCVDIDECSIMNGGCETFCTNSEGSYECSCQPGFALMPDQR
                                                                                                                                                      MGNFFCLCKAGWGG----RLCDKDVNECSQENGGCLQICHNKPGSFHCSCHSGFELSSDGR 193
                                                                                                                                                                                       CLNTEGS-----YRC-----ECPPGHQLAPNISACIDINECELSAHLCPHGRCVNL
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ision 20-Feb-1995 #text_change
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Result
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              88
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1: uniprot_sprot:*
2: uniprot_trembl:*
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3675
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  GenCore version 5.1.9 (c) 1993 - 2006 Biocceleration
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           PROS_HŪMAN
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Q16539 HUMAN
Q9NSDO_HUMAN
Q9NSDO_HUMAN
PROS_BOVIN
Q29054 PIG
PROS_RABIT
PROS_MOUSE
Q3TR66 MOUSE
Q3TR66 MOUSE
Q4SH13 TETNG
Q68E12 HUMAN
Q62628 RAT
Q68E19 MOUSE
Q4SHN1 TETNG
Q54YPO DICDI
Q7PPCO_ANOGA
FBLN1 MOUSE
Q3TWK8 MOUSE
Q3TWK8 MOUSE
Q3TWK8 MOUSE
Q3TWK8 MOUSE
                                                                                                                                                                                                                                                           Q6PAE0_XENLA
Q4SHY2_TETNG
Q7T3H4_BRARE
                                                                                                                                                                                                                                                                                             Q6IRI1 RAT
GAS6 MOUSE
Q2MJP6 NOTVI
Q6DFA5 XENLA
                                                                                                                                                                                                                                                                                                                                              Q6IMN1 HUMAN
GAS6 HUMAN
GAS6 RAT
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                                                                                                                     Q3tr66
P53813
Q4shy3
                                                                                                                                                                                                                                                                                           Q2mjp6
Q6dfa5
           Q3twk8
Q9vs89
                                                                                             Q65812
Q62628
                                                                                                                                                                                                                                                           Q4shy2
Q7t3h4
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Q14393
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Pfam;
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                                                                                                                                                                                                                                                                                                                                                         Matthews L
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| | ALIGNMENTS | | | | | |
|--------------------|--------------|----|------|-----|-------|----|
| Q75n88 homo sapien | Q75N88_HUMAN | N | 1365 | 9.6 | 352 | 5 |
| O88840 mus musculu | O88840_MOUSE | N | 3857 | 9.6 | 353 | 4 |
| 075413 homo sapien | 075413_HUMAN | N | 1624 | 9.6 | 353.5 | ü |
| 075412 homo sapien | 075412_HUMAN | N | 1557 | 9.6 | 353.5 | 2 |
| Q4su37 tetraodon n | Q4SU37_TETNG | N | 509 | 9.6 | 353.5 | Ë |
| | O00508 HUMAN | N | 1587 | 9.6 | 354.5 | ô |
| Q5ble3 brachydanio | Q5BLE3_BRARE | N | 469 | 9.7 | 356 | 39 |
| Q8k4g1 mus musculu | LTBP4 MOUSE | μ. | 1666 | 9.7 | 356.5 | 8 |
| Q3ugul mus musculu | Q3UGU1_MOUSE | N | 1021 | 9.7 | 356.5 | 37 |
| Q5xj74 brachydanio | Q5XJ74_BRARE | N | 477 | 9.7 | 356.5 | 36 |
| P87363 gallus gall | P87363_CHICK | N | 708 | 9.7 | 357 | 5 |
| Q9eqc6 mus musculu | Q9EQC6_MOUSE | N | 961 | 9.7 | 357.5 | 4 |
| Q6nzl8 mus musculu | Q6NZL8 MOUSE | N | 1018 | 9.7 | 358 | 3 |
| Q9jjs0 mus musculu | Q9JJS0_MOUSE | N | 997 | 9.8 | 358.5 | 32 |
| | | | | | | |

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RESULT 1

OGINNI HUMAN

OGINNI;

OT 10-MAY-2005, integrated into UniProtKB/TFEMBL.

DT 10-MAY-2005, integrated into UniProtKB/TFEMBL.

DT 10-MAY-2005, sequence version 1.

DT 07-MAR-2006, entry version 1.

DT 07-MAR-2006, pequence version 1.

OR HOMO SADISHMERT (Human).

OR HOMO SADISHMERT (Human).

OR HOMO SADISHMERT (Human).

OR HOMO SADISHMERT (Human).

RN (1) FAXID-9606;

RN (1) FAXID-9606;

RN (1) FAXID-9606;

RN MUCLEOTIDE SEQUENCE.

RA WILLEOTIDE SEQUENCE.

RA MAINCA X. Samoy L. L., Ramitez-Lorca R., Villar J., de Frutos P.G., Asala N.;

RI HUMAN vitamin K-dependent GAS6: gene structure, allelic variation, and association with stroke.";

RI HUMAN MUCLEOTIDE SEQUENCE.

RA SALA N.;

RI SUBMITTEE (MAY-2005) to the EMBL/GenBank/DDBJ databases.

CC EMBL/GenBank/DDBJ third parry amontation (TPA) entry

CC COPYIGHED YELD UNIPPOT CONSOUTIUM, see http://www.uniprot.org/terms

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Matches 678;
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SMART; SN
SMART; SN
PROSITE;
PROSITE;
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PROSITE;
PROSITE;
PROSITE;
GAS6 HUMAN
Q14393; Q7Z7N3;
07-JUN-2004, in
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T; SM00179; EGF CA; 3.
T; SM00069; GLA; 1.
T; SM00282; LamG; 2.
ITE; PS00010; ASX_HYDROXYL; 4.
ITE; PS000186; EGF 1; UNKNOWN_1.
ITE; PS01186; EGF 2; 3.
ITE; PS01187; EGF CA; 3.
ITE; PS01187; EGF CA; 3.
ITE; PS0001; GLA; I UNKNOWN_1.
ITE; PS0001; GLA; I UNKNOWN_1.
ITE; PS0005; LAM_G_DOMAIN; 2.
ENCE 678 AA; 74925 MW; BB6D8A
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                                                                                                                                                                                          DYHSTKKLKKQLVVLAVEHTALALMEIKVCDGQEHVVTVSLKDGEATLEVDGTRGQSEVS
                                                                                                                                                                                                                                                                                                           LTVGGIPFHEKDLVQPINPRLDGCMRSWNWLNGEDTTIQETVKVNTRMQCFSVTERGSFY
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                                                                                      AAQLQERLAVLERHLRSPVLTFAGGLPDVPVTSAPVTAFYRGCMTLEVNRRLLDLDEAAY
                                                                                                                                                                                                          DYHSTKKLKKQLVVLAVEHTALALMEI KVCDGQEHVVTVSLRDGEATLEVDGTRGQSEVS
                                                                                                                                                                                                                                                                     PGSGFAFYSLDYMRTPLDVGTESTWEVEVVAHIRPAADTGVLFALWAPDLRAVPLSVALV
                                                                                                                                                                                                                                                                                                                                                              YNGVGRVTSSGPVINHGMWQTISVEELARNLVIKVNRDAVMKIAVAGDLFQPERGLYHLN
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                                                                                                                                         AAQLQERLAVLERHLRSPVLTFAGGLPDVPVTSAPVTAFYRGCMTLEVNRRLLDLDEAAY
                                                                                                                                                                                                                                                                                             LTVGGIPFHEKDLVQPINPRLDGCMRSWNWLNGEDTTIQETVKVNTRMQCFSVTERGSFY
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UniProtKB/Swiss-Prot.
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Pred. No. 3.2e-250;
                         PRT;
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RA PLINMEG=14/02/39; DOI=10.1036/HJL-25;
RA OLa T. Suzuki Y., Nishikawa T., Otsuki T., Sugiyama T., Irie R.,
RA Wakamatsu A., Hayashi K., Sato H., Nagai K., Kimura K., Makita H.,
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Yoshikawa Y., Matsunawa H., Ichihara T., Shiohata N., Sano S.,
RA Moriya S., Momiyama H., Satoh N., Takami S., Terashima Y., Suzuki O.,
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RA Rawabata A., Hikiji T., Noguchi S., Itoh T., Shigeta K., Senba T.,
RA Mizushima-Sugano J., Satoh T., Shirai Y., Makashim Y., Nakagawa K.,
RA Mizushima-Sugano J., Satoh T., Shirai Y., Masuho Y., Yamasshita R.,
RA Nakai K., Yada T., Nakamura Y., Ohara O., Isogai T., Sugano S.;
RA Nakai K., Yada T., Nakamura Y., Ohara O., Isogai T., Sugano S.;
Ra Nakai K., Yada T., Nakamura Y., Ohara O., Isogai T., Sugano S.;
Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
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Rlausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
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Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
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Brownstein M.J., Uddin T.B., Toshiyuki S., Carninci P., Prange C.,
Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.G.
Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
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Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A
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Submitted
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                          TISSUE=Fetal lung, and Fetal spleen;
MEDLINE=2238857; PubMed=12477932; DOI=10.1073/pnas.242603899;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PubMed=14702039; DOI=10.1038/ng1285;
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Sumoy L., de Frutos P., Sala N.;
(MAR-2003) to the EMBL/GenBank/DDBJ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (ISOFORM
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                                                 wullahy S.J., ratne P.H., Hulyk S.W.,
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.c N.K.,
•h F.,
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x-RAY CRYSTALLOGRAPHY (2.2 ANGSTROMS) OF 261-721 (ISOFORM MUTAGENESIS OF PHE-530; LEU-663 AND TYR-703.)

4 PubMed=8621659; DOI=10.1074/jbc.271.16.9785;

5 Mark M.R., Chen J., Hammonds R.G., Sadick M., Godowski P.J.

7 "Characterization of Gas6, a member of the superfamily of containing proteins, as a ligand for Rse and Axl.";

1 Biol. Chem. 271:9785-9789(1996)

2 -i- FUNCTION: Ligand for tyrosine-protein kinase receptors and MER whose signaling is implicated in cell growth a cell adhesion and cell migration. Plays a role in that camplifying platelet aggregation and secretion in response aggoniates (By similarity).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RECEPTOR INTERACTION, SUBCELLULAR LOCATION, AND PROCESSING PubMede-9326369; DOI=10.1016/S0014-5793(97)0.1093-4; Goruppi S., Yamane H., Marcandalli P., Garcia A., Clogston Gostissa M., Varnum B., Schneider C.; "The product of a gas6 splice variant allows the release of responsible for Ax1 tyrosine kinase receptor activation."; FEBS Lett. 415:59-63(1997).
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PubMed=8939948; DOI=10.1074/jbc.271.47.30022;
Nagata K., Ohashi K., Nakano T., Arita H., Zong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     common ligand :
J. Biol. Chem.
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"Generation and initial." ""

"Generation and initial." ""
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Identification of the product of growth arrest-specific gene common ligand for Axl, Sky, and Mer receptor tyrosine kinases. J. Biol. Chem. 271:30022-30027(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ALTERNATIVE SPLICING (ISOFORM 1), AND TISSUE SPECIFICITY. PubMed=9326368; DOI=10.1016/S0014-5793(97)01094-6; Marcandalli P., Gostissa M., Varnum B., Goruppi S., Schnei "Identification and tissue expression of a splice variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Axl receptor tyrosine kinase stimulated by the vitamin K-dependent protein encoded by growth-arrest-specific gene 6."; Nature 373:623-626(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Varnum B.C., Young C., Elliott G., Garcia A., Bartley T.D., Fridell Y.W., Hunt R.W., Trail G., Clogston C., Toso R.J., Yanagihara D., Bennett L., Sylber M., Merewether L.A., Tsen Escobar E., Liu E.T., Yamane H.K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nagata K.,
Mizuno K.;
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                                                        SUBUNIT: Monomer (Probable)
SUBCELLULAR LOCATION: Secre
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Event=Alternative splicing;
                                ALTERNATIVE PRODUCTS:
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ett. 415:56-58(1997).
                                                           Secreted
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     Named isoforms=3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MIM; 600441; gene.
Reactome; Q14393;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ensemb1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PTM: Isoform 1 is proteolytically processed after secretion to yield a N-terminal 36 kDa protein and a C-terminal 50 kDa protein and a C-terminal 50 kDa protein contains which activates AXI. PTM: Gamma-carboxyglutamate residues are formed by vitamin K dependent carboxylation. These residues are essential for the binding of calcium (By similarity).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              binding of calcium (By similarity).
SIMILARITY: Contains 4 EGF-like domains.
SIMILARITY: Contains 1 Gla (gamma-carboxy-glutamate)
SIMILARITY: Contains 2 laminin G-like domains.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TISSUE SPECIFICITY: Plasma.
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                                                                                                                                                  121
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AY170372;
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                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                TPNPCDRKGTQACQDLMGNFFCLCKAGWGGRLCDKDVNECSQENGGCLQICHNKPGSFHC
ECLQGRCEQVCVNSPGSYTCHCDGRGGLKLSQDMDTCE
                                 SCHSGFELSSDGRTCQDIDECADSEACGEARCKNLPGSYSCLCDEGFAYSSQEKACRDVD
                                                           SCHSGFELSSDGRTCQDIDECADSEACGEARCKNLPGSYSCLCDEGFAYSSQEKACRDVD
                                                                                                          TPNPCDRKGTQACQDLMGNFFCLCKAGWGGRLCDKDVNECSQENGGCLQICHNKPGSFHC
                                                                                                                                                                                 LERECVEELCSREEAREVFENDPETDYFYPRYLDCINKYGSPYTKNSGFATCVQNLPDQC
                                                                                                                                                                                                           LERECVEELCSREEAREVFENDPETDYFYPRYLDCINKYGSPYTKNSGFATCVQNLPDQC
                                                                                                                                                                                                                                                              MAPSISPGPAALRRAPQIIIIIIIAAECALAALIPAREATQFIRPRQRRAFQVFEEAKQGH
                                                                                                                                                                                                                                                                                        MAPSLSPGPAALRRAPQLLLLLLLAAECALAALLPAREATQFLRPRQRRAFQVFEEAKQGH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ENSG00000183087;
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X-ray; A/B=322-721.
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AAH38984.
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AAO84057
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94.0%;
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ental confirmation available;
Y: Plasma. Isoform 1 and isoform
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GAS6 RAT
Q63772;
07-JUN-2004,
Nagata K., Ohashi
Mizuno K.;
"Identification
                                                                                                                                                                                                                                                                                                                               01-NOV-1996, sequence version 1
07-MAR-2006, entry version 46.
Growth-arrest-specific protein
potentiating factor) (GPP).
Name=Gas6;
                                                                                                                                                                                                                                                         Mammalia; Eutheria; Euarchor
Muroidea; Muridae; Murinae;
NCBI_TaxID=10116;
                                                                                                                                                                                                                                                                                                                                                                                                                                 RAT
                                                                                                           RECEPTOR INTERACTION,
PubMed=7559388; DOI=10
Ohashi K., Nagata K.,
Suzuki K., Mizuno K.;
                      RECEPTOR INTERACTION.
PubMed=8939948; DOI=10.1074/jbc.271.47.30022;
Nagata K., Ohashi K., Nakano T., Arita H., Zo
                                                                    Suzuki K., Mizuno K.;
"Stimulation of sky receptor tyrosine arrest-specific gene 6.";
J. Biol. Chem. 270:22681-22684(1995).
                                                                                                                                                                                                           Ohara O.,
                                                                                                                                                                                  potentiating
                                                                                                                                                                                                                       Nakano
                                                                                                                                                                                                                                  NUCLEOTIDE SEQUENCE
MEDLINE=95197586; Pu
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                                                                                                                                                                                               "Vascular
                                                                                                                                                                                                                                                                                                                      Rattus norvegicus (Rat).
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                                                                                                                                                                    smooth muscle cell-derived, Gla-containing growth-
ing factor for Ca(2+)-mobilizing growth factors.";
Chem. 270:5702-5705(1995).
                                                                                                                                                                                                           Arita H.;
                                                                                                                                                                                                                                                                                             ; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
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                                                                                                                                   DOI=10.
 of the product
                                                                                                                                                                                                         CE [MRNA].
PubMed=7890695; DOI=10.1074/jbc.
PubMed=784061 N., Kishino J., !
                                                                                                                     SUBCELLULAR LOCATION, A 0.1074/jbc.270.39.22681; Toshima J., Nakano T.,
                                                                                                                                                                                                                                                                                    Rattus.
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InterPro; IPR000294; VitK_de
Pfam; PP00008; EGF: CA; 2.
Pfam; PF07645; EGF CA; 2.
Pfam; PF002504; Gla: 1.
Pfam; PF002504; Gla: 1.
Pfam; PF002501; Laminin G_2;
PRINTS; PR00001; GLABLGOD.
SMART; SM00181; EGF; 4.
SMART; SM0019; EGF; CA; 3.
SMART; SM00069; GLA; 1.
SMART; SM00089; LAmG; 2.
SMART; SM00089; LAmG; 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              common ligand f
J. Biol. Chem.
[4]
                                                                                             PROSITE; PS00010; A
PROSITE; PS00022; E
PROSITE; PS01186; E
PROSITE; PS50026; E
PROSITE; PS001187; E
PROSITE; PS00011; G
PROSITE; PS50998; G
PROSITE; PS50998; G
Calcium; EGF-like d
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Arita H.;

activity of Gas6: contribution of endogenous Gas6 to the proliferation of vascular smooth muscle cells.";

Biochem. J. 323:387-392(1997).
DOMAIN
DOMAIN
DOMAIN
DOMAIN
DOMAIN
                                                               Growth
SIGNAL
CHAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               amplifying platelet aggregation and secretion in respon agonists (By similarity).

SUBUNIT: Monomer (By similarity).

SUBCELLULAR LOCATION: Secreted protein.

TISSUE SPECIFICITY: Plasma.

PIM: Gamma-carboxyglutamate residues are formed by vital dependent carboxylation. These residues are essential fibinding of calcium (Probable).

SIMILARITY: Contains 1 GGla (gamma-carboxy-glutamate) do similarity: Contains 2 laminin G-like domains.
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InterPro;
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InterPro;
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Q63772; 258-674.
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                                                                                     regulation;
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 50
113
153
194
235
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                                                                                             ; ASX HYDROXYL; 4
; EGF 1; 1.
; EGF 2; 3.
; EGF 3; 4.
; EGF CA; 3.
; GLA 1; 1.
; GLA 1; 1.
; GLA 1; 1.
; GLA 2; 1.
; LAM G DOMAIN; 2
e domain; Gamma-
                                                             uomain; Gamma-carboxyglutamic acid;
Metal-binding; Repeat; Signal.
27 Potential.
674
  91
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275
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  /FTId=PRO_Gla.

EGF-like 1
EGF-like 2
EGF-like 3
EGF-like 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mRNA.
                                                      Growth-arrest-specific /FTId=PRO_0000007591.
                                                                                                                                                                                                                                                                                                                                                                                                                                      norvegicus
 e 1; calcium-binding
e 2; calcium-binding
e 3; calcium-binding
e 4; calcium-binding
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CARBOHYD
CARBOHYD
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DISULFID

661 665

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Similarity
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                                                                                                                                                                                                                                                                                                                                                               DRKGTQACQDLMGNFFCLCKAGWGGRLCDKDVNECSQENGGCLQICHNKPGSFHCSCHSG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              VEELCSREEAREVFENDPETDYFYPRYLDCINKYGSPYTKNSGFATCVQNLPDQCTPNPC
                    KKLKKQLVVLAVEHTALALMEIKVCDGQEHVVTVSLRDGEATLEVDGTRGQSEVSAAQLQ
                                                                                            AFYSLDYMRTPLDVGTESTWEV
                                                                                                                               IPFHEKDLVQPINPRLDGCMRSWNWLNGEDTTIQETVKVNTRMQCFSVTERGSFYPGSGF
                                                                                                                                                                                         RITSSGPTINHGMWQTISVEELDRNLVIKVNKDAVMKIAVAGGLFQLERGLYHLNLTVGG
                                                                                                                                                                                                                                                 VIRLRFKRLQPTRLLAEFDFRTFDPEGVLFFAGGRSDSTWIVLGLRAGRLELQLRYNGVG
                                                                                                                                                                                                                                                                                                                                                                               FELSSDGRTCQDIDECADSEACGEARCKNLPGSYSCLCDEGFAYSSQEKACRDVDECLQG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PGP-AALRRAPQLLLLLLAAECALAALLPAREATQFLRPRQRRAFQVFEEAKQGHLEREC
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82.9%;
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Calcium (By similarity).

N-linked (GLCNAc. . .) (Potential).

By similarity.

By similarity.
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Laminin G-like 2.
Calcium (By similarity)
Calcium (via carbonyl o similarity).
Calcium (via carbonyl c Calcium (via carbonyl c
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Pred.
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ced. No. 7.3e
Mismatches
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RX Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., McKernan K.J., Abramson R.D., Mullahy S.J.,
RA Brownstein M.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Bukesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
"Generation and initial analysis of more than 15,000 full-length human roll mouse cDNA sequences.";

Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
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QGIRL1;
QGIRL1;
05-JUL-2004,
05-JUL-2004,
07-FEB-2006,
Gas6 protein.
Name=Gas6;
    InterPro;
                                                                                                                              SMR; Q6IRLI; 258-674.
Ensembl; ENSRNOG0000018233; Rattus nor G0; G0:0005509; C:cattracellular region; G0; G0:0005509; F:calcium ion binding; InterPro; IPR000152; Asx hydroxyl S.
InterPro; IPR013320; ConA_like_subgrp.
                                                                                                                                                                                                     EMBL;
HSSP;
SMR; Q
                                                                                                                                                                                                                                                            Copyrighted
Distributed
                                                                                                                                                                                                                                                                                                      Strausberg R.;
Submitted (MAY-2004)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TISSUE=Lung;
MEDLINE=22388257;
                                                                                                                                                                                                                                                                                                                                    TISSUE=Lung;
                                                                                                                                                                                                                                                                                                                                                 NUCLEOTIDE
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Mammalia; Eutheria;
Muroidea; Muridae; M
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P00736; 1
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IPR013320

IPR006210

IPR000742

IPR001881

IPR013091

IPR013032

IPR013032

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a; Chordata; Craniata; Vertebrata; Euteleostomi;

a; Eburchontoglires; Glires; Rodentia; Sciurognathi;

; Murinae; Rattus.
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EGF 2.

EGF Ca bd 2.

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ion; IEA.
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126 63 66 S 7 Query Match Best Local S Matches 558

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PÉAM; PF00594; GIA; 1.

PÉAM; PF00210; LAMÍNIN G 2; 2.

PEAM; PR00210; JEAMÍNIN G 2; 2.

PRINTS; PR00000; GLABLÃOD.

SMART; SM00181; EGF CA; 3.

SMART; SM00189; GLA; 1.

SMART; SM00109; GLA; 1.

SMART; SM0002010; ASX_HYDROXYL; 4.

PROSITE; PS00010; ASX_HYDROXYL; 4.

PROSITE; PS01106; EGF 2; 3.

PROSITE; PS01106; EGF 2; 3.

PROSITE; PS01187; EGF CA; 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     InterPro; IPR000294; VitK_d
Pfam; PF00008; EGF; 1.
Pfam; PF07645; BGF_CA; 2.
Pfam; PF00594; Gla; 1.
Pfam; PF02210; Laminin_G_2;
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                                                                                                               KKLKKQLVVLAVEHTALALMEIKVCDGQEHVVTVSLRDGEATLEVDGTRGQSEVSAAQLQ
                                                                                                                                                                                                                                                                                                         RVTSSGPVINHGMWQTISVEELARNLVIKVNRDAVMKIAVAGDLFQPERGLYHLNLTVGG
                                                                                                                                                                                                                                                                                                                                              VIRLREKRLQPTRLVABEDERTFDPEGILLFAGGHQDSTWIVLALRAGRLELQLRYNGVG
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                                                                                                                                                                         AFYSLNYTRTSLDVGTETTWEVEVVARIRPATDTGVLMAL-VGDKDVVLLSVALVDYHST
                                                                                                                                                                                       AFYSLDYMRTPLDVGTESTWEVEVVAHIRPAADTGVLFALMAPDLRAVPLSVALVDYHST
                                                                                                                                                                                                                                 IPFKESDLVQPINPRLDGCMRSWNWLNGEDSAIQETVKANTKMQCFSVTERGSFFPGNGF
                                                                                                                                                                                                                                                IPFHEKDLVQPINPRLDGCMRSWNWLNGEDTTIQETVKVNTRMQCFSVTERGSFYPGSGF
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TSHSCPPVEHVTA
                         TAHSCPPVEPAAA
                                                                                  ERLAVLERHLRSPVLTFAGGLPDVPVTSAPVTAFYRGCMTLEVNRRLLDLDEAAYKHSDI
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 674
                           678
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                                                          /GGLPDVQVTSTPVTAFYRGCMTLEVNGKTLDLDTASYKHSDI
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Pred. No. 1
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RESULT 5
GASG MOUS
GASG MOUS
GASG MOUS
GASG MOUS
AC Q615
DT Q7-Q
DT Q7
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RA KIRUSHORE-2238557; Pubmeda:1247932; DOI=10.1073/pnas.242603899;
RA KIRUSHOR R.D., Collins F.S., Wagner L.H., Derge J.G.,
RA KIRUSHOR R.D., Collins F.S., Wagner L., Shaefer C.F., Bhat N.K.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., McKernan K.J., Malek J.A., Gunaratrue P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Richards S., Wadan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Rhiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
RA "Generation and initial analysis of more than 15,000 full-length human
"Generation and initial analysis of more than 15,000 full-length human
"Tand mouse cDNA sequences";
Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
                               Copyrighted
Distributed
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Q61592; Q99K57;
Q7-JUN-2004, integrated into UniProtKB/Swiss-Prot.
Q7-JUN-2004, sequence version 2.
Q7-MAR-2006, entry version 45.
Growth-arrest-specific protein 6 precursor (GAS-6)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PubMed=11175853; DOI=10.1038/84667;
Angelillo-Scherrer A. de Frutos P., Aparici
Lupu F., Arnout J., Dewerchin M., Hoylaerts
Dahlback B., Carmeliet P.;
"Deficiency or inhibition of Gas6 causes pla
protects mice against thrombosis.";
Nat. Med. 7:215-221(2001).
-!- FUNCTION: Ligand for tyrosine-protein ki
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MEDLINE=93330291; PubMed=8336730;
Manfioletti G., Brancolini C., Av
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Mammalia; Eutheria; Euarchontoglires;
Muroidea; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mus musculus (Mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FUNCTION
                                                                                                                        response to known agonists.
SUBUNIT: Monomer (By similarity).
SUBCELLULAR LOCATION: Secreted protein (By similarity).
PTM: Gamma-carboxyglutamate residues are formed by vitamin dependent carboxylation. These residues are essential for the carboxylation of the similarity).
MISCELLANEOUS: GAS6 deficient mice show protection against thrombosis, but no spontaneous bleeding.
SIMILARITY: Contains 4 EGF-like domains.
SIMILARITY: Contains 1 Gla (gamma-carboxy-glutamate) domain SIMILARITY: Contains 2 laminin G-like domains.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and MER whose signaling is implicated in cell cell adhesion and cell migration (By similarit thrombosis by amplifying platelet aggregation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FUNCTION: Ligand for tyrosine-protein and MER whose signaling is implicated
                                  l by the I
                                     e UniProt Consortium, see http://www.uniprot.org/terms
the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Craniata; Vertebrata; Euteleostomi;
oglires; Glires; Rodentia; Sciurogna
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ylaerts M., Herbert
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Collen
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Ensembl; ENSMSGOODOO31451; Mus muscullus MGI; MGI; 95660; Gas6.

GO: GO:0005615; C:extracellular space; T; InterPro; IPR000152; Asx hydroxyl S. InterPro; IPR000152; Asx hydroxyl S. InterPro; IPR000152; Asx hydroxyl S. InterPro; IPR0001320; Con like subgrp. InterPro; IPR000742; EGF 3.

InterPro; IPR000742; EGF 3.

InterPro; IPR001881; EGF Ca bd. 2.

InterPro; IPR001891; EGF Tike reg. InterPro; IPR0013032; EGF Tike reg. InterPro; IPR0013032; EGF Like reg. InterPro; IPR0013032; EGF Like reg. InterPro; IPR001383; GLA blood. InterPro; IPR001294; VitK dep_GLA. InterPro; IPR001294; VitK dep_GLA. Pfam; PF000081; EGF; 1.

Pfam; PF000081; EGF; 4.

Pfam; PF02210; Laminin G 2; 2.

PRINTS; SM00101; GLABLOOD.

SMART; SM00101; EGF; 4.

SMART; SM00102; EGF 1; 1.

PROSITE; PS00101; ASX HYDROXYL; 4.

PROSITE; PS00102; EGF 2; 3.

PROSITE; PS00118; EGF 2; 3.
      METAL
CARBOHYD
CARBOHYD
DISULFID
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EMBL; BC00544
PIR; A48089;
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BC005444;
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4; AAH05444.1;
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Metal-binding; Repeat; Signal.
27 Potential.
674
      652
417
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1130
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1177
1177
1192
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233
2488
274
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Growth-arrest-specific protein 6.
/FTId=PRO_000007590.
Gla.
BGF-like 1; calcium-binding (Potenti.
BGF-like 2; calcium-binding (Potenti.
EGF-like 3; calcium-binding (Potenti.
EGF-like 4; calcium-binding (Potenti.
EGF-like 4; calcium-binding (Potenti.
Laminin G-like 1.
Laminin G-like 2.
Calcium (By similarity).
Calcium (via carbonyl oxygen) (By
similarity).
Calcium (via carbonyl oxygen) (By
similarity).
N-linked (GlcNAc. . .) (Potential).
By similarity.
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Best Local
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RESULT 6
Q2MJP6 NOTVI
Q2MJP6 NOTVI
ID Q2MJP6;
AC Q2MJP6;
DT 07-FEB-2006, integrated into Uni
DT 07-FEB-2006, sequence version 1.
DT 07-FEB-2006, entry version 1.
DE Growth arrest-specific 6.
GN Name-Gas6;
OS Notophthalmus viridescens (Easte OC Eukaryota; Metazoa; Chordata; Cr OC Amphibia; Batrachia; Caudata; Sa OC Notophthalmus.
OX NCBI_TaxID-8316;
RN [1]
RP NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                        AHSCPPVEPA
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48; Conservative
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                                                                        viridescens (Eastern newt) (Tri
tazoa; Chordata; Craniata; Vertel
rachia; Caudata; Salamandroidea;
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Pred. No. 7.3e-203;
5; Mismatches 75;
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7C41F7693903F401
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   C.;
                                                                         ) (Triturus viridescens).
Vertebrata; Euteleostomi;
Didea; Salamandridae;
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                                       16-AUG-2004, integrated into UniProtKB/TrEMBL 16-AUG-2004, sequence version 1. 07-FEB-2006, entry version 13.
           Gas6-prov protein.
Name=gas6-prov;
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W; 9D860E80336BAD64
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RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Biatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gy L.J., Hulyk S.W.,
RA Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Glbbs R.A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
RA Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;
RT "Generation and initial analysis of more than 15,000 full-length human
RT and mouse cDNA sequences.";
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    PROSITE; PS00010; ASX_HYDROXYL; PROSITE; PS00022; EGF_1; 1. PROSITE; PS01186; EGF_2; 3.
                                                                                                                                                                                                                                                                                                                                       GO; GO:0005576; C:extracellular region GO; GO:0005509; F:calcium ion binding;
                                                                                                                                                                                                                                                                                                                                                                                                Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (JUL-2004) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                       TISSUE=Oocytes;
Klein S., Gerha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=22341132; PubMed=12454917; DOI=10.1002/dvdy.10174;
Klein S.L., Strausberg R.L., Wagner L., Pontius J., Clifto
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                                            ; PF02210; Laminin G 2; TS; PR00001; GLABLOOD. TS; SM00181; EGF; 4. T; SM00179; EGF CA; 3. T; SM00069; GLA; 1. T; SM00082; LamG; 2.
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IPR013091;
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IPR001791;
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IPR013320; ConA_like_su
IPR006210; EGF.
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EGF Ca bd.
EGF Ca bd_2.
EGF like.
EGF like reg.
GLA blood.
                                                                                                                                                                     Laminin_G.
Laminin_G 2.
VitK_dep_GLA.
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Clifton S.W., NIH Xenopus

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Name=MGC8463 protein.

Name=MGC8463 protein.

Name=MGC8463; Carcian clawed frog).
Ekkaryota; Metazoa; Chordata; Craniata; Vertebr; Ekkaryota; Metazoa; Chordata; Mesobatrachia; Pipo:
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PROSITE;
PROSITE;
SEQUENCE
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IKRQFLILSIENIVVSRIEVQVCD-SEHIVEISASINDLFLSFDGTLGQKELPESQMRTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PS50026;
PS01187;
PS00011;
PS50098;
PS50025;
P669 AA;
                                                                                                                                                                                                                                                         HSCPPIK
                                                                                                                                                       HSCPPVE
                                                                                                                                                                                                                                                  PHIDY
                                                                                                                                                                                                                                                                                              FHEKDLVQPINPRLDGCMRSWNWLNGEDTTIQETVKVNTRMQCFSVTERGSFYPGSGFAF
                                                                                                                                                                                                                                                                                                                                 TSSGPVINHGMWQTISVEELARNLVIKVNRDAVMKIAVAGDLFQPERGLYHLNLTVGGIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FKSSELTQSINPRLDGCMRNWNWLNGEDTSIQETVRMSEKNQCFLVAGRGSFFPGRGFAL
                                                                                                                                                                                                                                                                                                                          TSSGPLINHGMWQTIS'
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                                                                                                                                                                                                                                                                                                                                                                                                        EQVCVNSPGSYTCHCDGRGGLKLSQDMDTCEDILPCVPFSVAKSVKSLYLGRMFSGTPVI
                                                                                                                                                                                                                                                                                                                                                                                                                                             PSALTATWILVTLLIAEPLHGTILLSAKDASQFLRQRQRRANQIFEETKQGHLERECVEE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PAALRRAPQLLLLLLAAECALAALLPAREATQFLRPRQRRAFQVFEEAKQGHLERECVEE
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Xenopus;
                                                                                                                                                                                                                                               -TYPSKEETEANWRVEVNAQINPATDTGVLFALVSKDM-VVPLSLALIDYHSSAK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; EGF_3; 4.
; EGF_CA; 2.
; GLA_1; UNKNOWN_1.
; GLA_2; 1.
; LAM_G DOMAIN; 1.
; AM_G DOMAIN; ABDE:
                                                                                                                                        999
                                                                                                                                                        674
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             62.4%;
Xenopus
                                                                                                                                                                                                                                                                                                                         VEELDRNLVIKVNKDAVMKIAASGDLFSVDKGLYQLNLTVGGIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     105;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 2293.5;
Pred. No. 6.7e-
05; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A8DE1E362540F643 CRC64;
       Vertebrata; Euteleostomi; ia; Pipoidea; Pipidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7e-155;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    144;
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RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Stapleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Roberts S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Robak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Mhiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;
"Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences.";
Pfam; PF02210; Laminin G 2; 1.

PRINTS; PR00001; GLABLOOD.

SMART; SM00181; EGF; 4.

SMART; SM00179; EGF_CA; 3.

SMART; SM00069; GLA; 1.

SMART; SM00069; GLA; 1.

SMART; SM00282; LamG; 2.

PROSITE; PS0010; ASX HYDROXYL; 3.

PROSITE; PS01186; EGF_2; 3.

PROSITE; PS01187; EGF_CA; 2.
                                                                                                                                                                                                          HSSP; P00743; 1APO.

GO; GO:0005576; C:extracellular region; II
GO; GO:0005576; C:extracellular region; II
GO; GO:0005509; F:calcium ion binding; IE;
InterPro; IPR000152; Asx hydroxyl S.
InterPro; IPR000320; ConĀ_like_subgrp.
InterPro; IPR000742; EGF 3.
InterPro; IPR000742; EGF 3.
InterPro; IPR0001881; EGF Ca bd 2.
InterPro; IPR001881; EGF Ca bd 2.
InterPro; IPR0110032; EGF like_reg.
InterPro; IPR013032; EGF like_reg.
InterPro; IPR013032; EGF like_reg.
InterPro; IPR001791; Laminin_G.
InterPro; IPR012680; Laminin_G.2.
InterPro; IPR000294; VitK_dep_GLA.
Pfam; PF00008; EGF; 1.
Pfam; PF000784; EGF, CA; 3.
Pfam; PF007645; EGF_CA; 3.
                                                                                                                                                                                                  Pfam;
Pfam;
Pfam;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Copyrighted by the UniProt Consortium, see http://www.uniprot.org/tuDistributed under the Creative Commons Attribution-NoDerivs License
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Submitted (OCT-2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TISSUE=Liver;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUCLEOTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Richardson P.;
"Genetic and genomic initiative.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=22341132; PubMed=12454917; DOI=10.1002/dvdy.10174; Klein S.L., Strausberg R.L., Wagner L., Pontius J., Clifton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TISSUE=Liver;
MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     initiative.
                                                                                                                                                                                                PF00594;
PF02210;
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RESULT
Q4SHY2
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DT 19
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PROSITE;
PROSITE;
                                                                                                                    Tetraodon nigroviridis (Green puffer).

Bukaryota; Metazoa; Chordata; Cranitata; Vertebrata; Euteleostomi;

Actinopterygii; Neopterygii; Teleostei; Buteleostei; Neoteleostei;

Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontiformes;

Tetradontoidea; Tetraodontidae; Tetraodon.
                                                                                                                                                                                                                                        19-JUL-2005, integrated into UniProtK
19-JUL-2005, sequence version 1.
07-FEB-2006, entry version 7.
Chromosome 5 SCAF14581, whole genome
ORFNames=GSTENG00017935001;
                                                                                                                                                                                                                                                                                                                                                            Q4SHY2_TETNG
Q4SHY2;
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NUCLEOTIDE SEQUENCE.
PubMed=15496914; DOI=10.1038/nature03025;
Jaillon O., Aury J.-M., Brunet F., Petit
                                                                                                                                                                                                                                                                                                                                                                                                           TETNG
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PS50998; C
PS50025; I
668 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PINPRLDGCMRSWNWLNGEDTTIQETVKVNTRMQCFSVTERGSFYPGSGFAFYSLDYMRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HGMWQTISVEELARNLVIKVNRDAVMKIAVAGDLFQPERGLYHLNLTVGGIPFHEKDLVQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DIDECAESPNICGTAQCKNVRSTYVCLCEDGYRYDELTKSCQDVDECKEGRCEQTCVNSP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DIDECADS-EACGEARCKNLPGSYSCLCDEGFAYSSQEKACRDVDECLQGRCEQVCVNSP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HGDFFCHCKPGWTGKSCGTDINECATENGNCSQICINKPGTHQCLCSSGYRLHANNKLCI
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                                                                                                                                                                                                                                                                                                                                                                                    PRELIMINARY;
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GLA_2; 1.
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Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                     UniProtKB/TrEMBL
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       Petit J.-L.,
                                                                                                                                                                                                                                                                shotgun sequence.
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    Stange-Thomann
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Nicaud S., Jaffe D., Fisher S., Lutfalla G., Dossat C., Segurens B.,
A Dasilva C., Salanoubat M., Levy M., Boudet N., Castellano S.,
Anthouard V., Jubin C., Castelli V., Katinka M., Vacherie B.,
A Anthouard V., Jubin C., Castelli V., Katinka M., Vacherie B.,
A Desilva C., Skalli Z., Cattolico L., Poulain J., De Berardinis V.,
A Cruaud C., Duprat S., Brottier P., Coutanneau J.-P., Gouzy J.,
A Cruaud C., Lardier G., Chapple C., McKernan K.J., McBean P., Bosak S.,
A Kellis M., Volff J.-N., Güigo R., Zody M.C., Mesirov J.,
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A Kellis M., Volff J.-N., Guigo R., Zody M.C., Mesirov J.,
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Best Local S
Matches 333
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Pfam; PF00294; Gla; 1.
Pfam; PF02210; Laminin G 2; 2.
PRINTS; PR00001; GLABLOOD.

SMART; SM00181; EGF; 4.
SMART; SM00181; EGF; 4.
SMART; SM00179; EGF_CA; 3.
SMART; SM00069; GLA; 1.
SMART; SM00069; GLA; 1.
SMART; SM000282; LamG; 2.
PROSITE; PS00010; ASX HYDROXYL; 4.
PROSITE; PS00022; EGF 1; 1.
PROSITE; PS01186; EGF 2; 3.
PROSITE; PS01186; EGF 2; 3.
PROSITE; PS01187; EGF_CA; 3.
PROSITE; PS01187; EGF_CA; 3.
PROSITE; PS01187; EGF_CA; 3.
PROSITE; PS05098; GLA—2; 1.
PROSITE; PS50998; GLA—2; 1.
PROSITE; PS50998; GLA—2; 1.
PROSITE; PS50998; GLA—2; 1.
PROSITE; PS50998; GLA—2; 1.
PROSITE; PS5098; GLA—2; 1.
PROSITE; PS50998; GLA—2; 1.
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GO; GO:0005509; F:calcium ion binding; IEE
InterPro; IPR000152; Asx hydroxyl_S.
InterPro; IPR006210; EGF.
InterPro; IPR000742; EGF_3.
InterPro; IPR001881; EGF_Ca bd.
InterPro; IPR013032; EGF_Like reg.
InterPro; IPR013032; EGF_Like reg.
InterPro; IPR001791; Laminin_G.
InterPro; IPR001791; Laminin_G.
InterPro; IPR000294; VitK dep_GLA.
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Genoscope; Whitehead Institute Centre for Genome Research;
Submitted (FEB-2004) to the EMBL/GenBank/DDBJ databases.
-!- CAUTION: The sequence shown here is derived from an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Copyrighted by the UniProt Consortium, Distributed under the Creative Commons
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EGF-like
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                                                                                                                        CDKDVNECSQENGGCLQICHNKPGSFHCSCHSGFELSSDGR-TCQDIDECADSEACGEAR
                                                                                                                                                                                                                                                                          LDCINKYGSPYTKNSGFATCVQNLPDQCTPNPCDRKGTQACQDLMGNFFCLCKAGWGGRL
                                                                                                                                                                                                                                                                                                                                                   CDTDVDECGKRNGGCDHRCNNTMGSYRCSCHQGYEL--HGRHTCADVDECKDPEVCGTAR
                                                                                                                                                                                                                   LACVERFGDAERKKODLITCVHNIPDOCSPSPCHPGGTVRCEDKKGDFLCHCFTGWAGAR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pred. No. 7.8e
2; Mismatches
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Attribution-NoDerivs License
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RA MEDILINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Willalon D.K., Wuzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Schnern S.Schein J.E., Jones S.J.M., Marra M.A.;
RA Schnerach A., Schein J.E., Jones S.J.M., Marra M.A.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         O7T3H4_BRARE PRELIMINARY; PRT; 648 AA. Q773H4; 01-OCT-2003, integrated into UniProtKB/TrEMBL. 01-OCT-2003, sequence version 1. 07-FEB-2006, entry version 15.
NUCLEOTIDE SEQUENCE TISSUE=Embryo;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Brachydanio rerio (Zebrafish) (Danio rer
Eukaryota; Metazoa; Chordata; Craniata;
Actinopterygii; Neopterygii; Teleostei;
Cyprinidae; Danio.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORFNames=zgc:63860;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Growth arrest specific
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Ostariophysi; Cypriniformes;
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SMART; SM00082; LamG; 2.

PROSITE; PS00010; ASX HYDROXYL; 3.

PROSITE; PS00012; EGF 1; 1.

R PROSITE; PS01186; EGF 2; 3.

R PROSITE; PS50026; EGF 3; 3.

R PROSITE; PS500187; EGF CA; 2.

R PROSITE; PS500187; EGF CA; 2.

PROSITE; PS00011; GLA 1; UNKNOWN 1.

PROSITE; PS00011; GLA 2; 1.

PROSITE; PS50025; LAM G DOMAIN; 2.

PROSITE; PS00228; TUBŪLĪN B AUTOREG;

CROTIENCE 648 AA; 72626 MW; E2935
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Best Local S
Matches 334
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ZFIN; ZDB-GENE-030131-7773;
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HSSP; Q14393; 11
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GO:0005576; C:extracellular region; IEA
GO:0005509; F:calcium ion binding; IEA.
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; SM00179; EGF CA; 2.
; SM00069; GLA; 1.
; SM00282; LamG; 2.
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                       RLVAEFDFRTFDPEGILLFAGGHQDSTWIVLALRAGRLELQLRYNGVGRVTSSGPVINHG
GFTAEFDLRTFDPEGVIFFAGGHLNSSWIVLLVHHGKLELQLKYGVVSRVTSSGPQVNDG
                                                                                         FRCFCDGF
                                                                                                              DECVETPDVCGSAHCSNLIGGLECLCDEGFIYDNISRSCVDVDECETHVCEEECVNTPGS
                                                                                                                                                                                                   DECADS-EACGEARCKNLPGSYSCLCDEGFAYSSQEKACRDVDECLQGRCEQVCVNSPGS
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; EGF_CA; 3.
; Gla; 1.
; Laminin_G_2;
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); ConA_like_subgrp.
EGF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; Laminin_G.
; Laminin_G 2.
; VitK_dep_GLA.
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EGF Ca bd.

EGF Ca bd 2.

EGF like.

EGF like reg.

GLA blood.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 47.8%; Score 1755; 50.8%; Pred. No. 1
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W; E2935F42AF21CF64
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                                                                                       IGLDRPLDMRRNSRSLYLGRMFSG:
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Attribution-NoDerivs
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                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=91084444; PubMed=2148110; Schmidel D.K., Tatro A.V., Phelps L. "Organization of the human protein S "Organization of the human protein S Biochemistry 29:7845-7852(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Euarchonto
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Name=PROS1; Synonyms=PROS;
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01-APR-1988, integrated into U
01-APR-1988, sequence version
07-MAR-2006, entry version 86.
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"SeattleSNPs. NHLBI HL66682 program for genom FHCRC, Seattle, WA (URL: http://pga.gs.washing Submitted (MAY-2003) to the EMBL/GenBank/DDBJ
                                                                    Rieder M.J., Carrington D.P., da Pont
Ahearn M.O., Kuldanek S.A., Rajkumar
Nickerson D.A.;
                                                                                                                                                                                                                                                     MEDLINE=91084445; PubMed=2148111;
Ploos van Amstel H.K., Reitsma P.H.,
"Intron-exon organization of the act
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=9606
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=88005138; PubMed=2820795; DOI=10.1016/0014-5793(87)80217-X; Ploos van Amstel H.K., van der Zanden A.L., Reitsma P.H.,
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                                                                                                                                                                                                                 evolution.";
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                                                                                  SEQUENCE [GENOMIC DNA].

Carrington D.P., da Ponte S.

Rajkumar N.,
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f the active human protein S gene PS a
duplication and silencing during prim
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RA Strausberg R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altechul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altechul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M.J., Ugdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Ugdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Raha S.S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Pahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Rodriguez A.C., Grinwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grinwood J., Schmutz J., Wyers R.M.,
RA Rodriguez A.C., Grinwood J., Schmutz J., Wyers R.M.,
RA Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;
"Generation and initial analysis of more than 15,000 full-length human
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                                                                                                                                                                                                                           Hayashi T., Nishioka J., Shigekiyo T., Saito S., Suzuki K.; "Protein S Tokushima: abnormal molecule with a substitution Lys-155 in the second epidermal growth factor-like domain of
                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=90335440; PubMed=2143091;
Bertina R.M., Ploos van Amstel H.K.,
Leemhuis M.P., Deutz-Terlouw P.P., Va
"Heerlen polymorphism of protein S, dimorphism of residue 460.";
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Liu T., Qian W.-J., Gritsenko M.A., Cam
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MEDLINE=86313649; PubMed=2944113;
                                                                           DOI=10.1002/(SICI)1098-1004(1999)14:1<30::AID-HUMU4>3
Espinosa-Parrilla Y., Morell M., Souto J.C., Tirado I.
Fontcuberta J. Estivill X., Sala N.;
"Protein S gene analysis reveals the presence of a cos
                                                                                                                                      PRO-501 AND MET-559.
MEDLINE=99374922; PubMed=10447256;
                                                                                                                                                                                                                                                                                                                                                                                  Blood
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hydrazide chemistry, and mass spectrometry.";
J. Proteome Res. 4:2070-2080(2005).
                               deficiency.";
Hum. Mutat. 14:30-39(1999)
                                                                                                                                                                                                                                                                        MEDLINE=94129009;
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tl. Acad. Sci.
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   Anticoagulant protein C in t
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Linkhub; P07225; -...

GO: GO:0005576; C:extracellular region; ]

GO: GO:0004866; F:endoopeptidase inhibito.

GO: GO:0004866; F:endoopeptidase inhibito.

GO: GO:0007596; P:blood coagulation; TAS

InterPro: IPR00152; Asx hydroxyl S.

InterPro: IPR00152; Asx hydroxyl S.

InterPro: IPR00152; ConA like_subgrp.

InterPro: IPR001320; ConA like_subgrp.

InterPro: IPR001301; EGF 3.

InterPro: IPR013091; EGF Ca_bd_2.

InterPro: IPR0013032; EGF like_

InterPro: IPR0013032; EGF like_

InterPro: IPR001303; GIA_blood.

InterPro: IPR001269; Laminin_G.

InterPro: IPR012690; Laminin_G.

InterPro: IPR0012690; VitK_dep_GIA.

InterPro: IPR000294; VitK_dep_GIA.
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Reactome; P07225; -.
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fibrinolysis.
- SUBCELLULAR LOCATION: Secreted protei
- TISSUE SPECIFICITY: Plasma.
- DISEASE: Defects in PROS1 are the ca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3L; M15036; AAA36479.1; -; mRNA.
3L; Y00692; CAA68687.1; -; mRNA.
3L; Y00692; CAA68688.1; AIT_SEQ; mRNA.
3L; M57853; AAA60357.1; -; Genomic_DNA.
3L; M578840; AAA60357.1; JOINED; Genomic_DNA.
3L; M578841; AAA60357.1; JOINED; Genomic_DNA.
3L; M578842; AAA60357.1; JOINED; Genomic_DNA.
3L; M578843; AAA60357.1; JOINED; Genomic_DNA.
3L; M57885; AAA60357.1; JOINED; Genomic_DNA.
3L; M57850; AAA60357.1; JOINED; Genomic_DNA.
3L; M57851; AAA60357.1; JOINED; Genomic_DNA.
3L; M57852; AAA60357.1; JOINED; Genomic_DNA.
3L; M36554; AAA60180.1; JOINED; Genomic_DNA.
3L; M36555; AAA60180.1; JOINED; Genomic_DNA.
3L; M36554; AAA60180.1; JOINED; Genomic_DNA.
3L; M36555; AAA60180.1; JOINED; Genomic_DNA.
3L; M36556; AAA60180.1; JOINED; Genomic_DNA.
3L; M36557; AAA60180.1; JOINED; Genomic_DNA.
3L; M36559; AAA60180.1; JOINED; Genomic_DNA.
3L
                                                                                                                                                                                                                                                                                                                                                                        GO:0005576; C:extracellular region; NA
GO:0004866; F:endopeptidase inhibitor
GO:0007596; P:blood coagulation; TAS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (PROS1 deficiency) [MIM:176880]. It is associated will increased risk to develop thrombotic disease (thrombotic SIMILARITY: Contains 4 BGF-like domains.

SIMILARITY: Contains 1 Gla (gamma-carboxy-glutamate)
SIMILARITY: Contains 2 laminin G-like domains.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A35610;
1Z6C; NM
    PF07645;
PF00594;
PF00054;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HGNC: 9456; PROS1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NMR;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    by the UniProt Consortium,
             ; EGF CA; 2.
; Gla; 1
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Laminin_G_1; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A=200-286
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Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                   activity;
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                                                            PROS_MACMU STANDARD; PRT; 649 AA.

Q28520;
Q28520;
Q1-NOV-1997, integrated into UniProtKB/Swiss-Prot.

O1-NOV-1997, sequence version 2.

O7-MAR-2006, entry version 56.

Vitamin K-dependent protein S precursor (Fragment).

Name=PROS1; Synonyms=PROS;

Macaca mulatta (Rhesus macaque).

Macaca mulatta (Rhesus macaque).

EUKaryota; Metazoa; Chordata; Craniata; Vertebrata; E
EUKaryota; Macaca; Eutheria; Euarchontoglires; Primates; Catar
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Best Local
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Cercopithecidae; Cercopithecinae;
NCBI_TaxID=9544;
[1]
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Pfam; PF02210; Laminin G;
PRINTS; PR00001; GLABLOOD
SMART; SM00181; EGF; 4.
SMART; SM00179; EGF_CA; 3.
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NDIRAHSCPSV
              SDITAHSCPPV
                                                                DLQRQLAVLDKAMKAKVATYLGGLEDVPFSATEVNAFYNGCMEVNINGVQLDLDEAISKH
                                                                                   QLQERLAVLERHLRSPVLTFAGGLPDVPVTSAPVTAFYRGCMTLEVNRRLLDLDEAAYKH
                                                                                                                                                           TKKLKKQLVVLAVEHTAL-ALMEIKVC-DGQEHVVTVSLRDGEATLEVDGTRGQSEVSAA
                                                                                                                                                                                                                                 FAFYSLDYMRTPLDVGTESTWEVEVVAHIRPAADTGVLFALWAPDLRAVPLSVALVDYHS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    VMLSNKKDCKDVDECSLKPSICGTAVCKNIPGDFECECPEGYRYNLKSKSCEDIDECSEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      YMSCKDGKASFTCTCKPGWQGEKCEFDINECKDPSNINGGCSQICDNTPGSYHCSCKNGF
                                                                                                                                        TSE-KSODILLSVENTVIYRIQALSLCSDQQSH--
                                                                                                                                                                                                            IAQFHIDYN----NVSSAEGWHVNVTLNIRPSTGTGVMLALVSGN-NTVPFAVSLVD--S
                                                                                                                                                                                                                                                                                                                  IPFH-EKDLVQPINPRLDGCMRSWNWLNGEDTTIQETVKVNTRMQCFSVTERGSFYPGSG
                                                                                                                                                                                                                                                                                                                                                     KITTGGDVINNGLWNMVSVEELEHSISIKIAKEAVMDINKPGPLFKPENGLLETKVYFAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RCEQVCVNSPGSYTCHCDGRGGLKLSQDMDTCEDILPCVPFSVAKSVKSLYLGRMFSGTP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               VFENDPETDYFYPRYLDCINKY-----GSPYTKNS--GFATCVQNLPDQCTPNPCDRKG
                                                                                                                                                                                                                                                                                 FPRKVESELIKPINPRLDGCIRSWNLMKQGASGIKEIIQEKQNKHCLVTVEKGSYYPGSG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TQACQDLMGNFFCLCKAGWGGRLCDKDVNEC---
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 676;
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Macaca

Vertebrata; Euteleostomi;

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PRESENTATION OF THE PROPERTY O
                                                                                                                                                                                                                                                                                                                                                                Pfam; PF00008; EGF; 1.

Pfam; PF000594; GGF; CA; 3.

Pfam; PF000594; Laminin G_1; 1

Pfam; PF000594; Laminin G_2; 2

PRINTS; PR00001; GLABLOOD.

SMART; SM00019; EGF; 4.

SMART; SM00019; GGA; 1.

SMART; SM00019; GGA; 1.

SMART; SM00028; LamG; 1.

SMART; SM00028; LamG; 1.

SMART; SM00028; EGF CA; 3.

SMART; SM00028; EGF 1; 1.

SMART; SM00028; EGF 2; 3.

PROSITE; PS00010; ASX HYDROXY PROSITE; PS01186; EGF 2; 3.

PROSITE; PS01186; EGF 3; 4.

PROSITE; PS01187; EGF CA; 3.

PROSITE; PS0001; GGA 1; 1.

PROSITE; PS0001; GGA 2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL; L31380;
HSSP; P00740;
SMR; Q28520;
InterPro; IPR
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InterPro;
InterPro;
InterPro;
Pfam; PF00
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MEDLINE=95134217; PubMed=7832752;
Greengard J.S., Fernandez J.A., Radtke
"Identification of candidate residues f
"ith C4b binding protein and activated
Biochem. J. 305;397-403(1995).
-!- FUNCTION: Anticoagulant plasma prot
DOMAIN
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SUBCELLULAR LOCATION: Secreted property of the secreted property of the secreted property.

TISSUE SPECIFICITY: Plasma.

SIMILARITY: Contains 4 EGP-like departs of the secretary.

SIMILARITY: Contains 2 laminin G-
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P00740;
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E; PSO0010; ASX HYDROXYL; 4.

E; PS00020; EGF-1; 1.

E; PS01186; EGF-2; 3.

E; PS01187; EGF-CA; 3.

E; PS01187; EGF-CA; 3.

E; PS0011; GLA-1; 1.

E; PS0001; GLA-2; 1.

E; PS00025; LAM-G DOMAIN; 2.

coagulation; Calcium; EGF-like domain.
carboxyglutamic acid; Glycoprotein; H;
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/FTIG=PRO 0000022121.
Vitamin K-dependent protein S
/FTIG=PRO 0000022122.
Gla.
EGF-like 1.
EGF-like 2; calcium-binding (
EGF-like 3; calcium-binding (
EGF-like 4; calcium-binding (
EGF-like 5; calcium-binding (
EGF-like 6; calcium-binding (
EGF-like 1; calcium-binding (
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N-linked (GlcNAc...
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Pfam; PF00054; Laminin G_1; Pfam; PF00054; Laminin G_2; Pfam; PF02210; GLAMININ G_2; PRINTS; PR00001; GLAMININ G_2; PRINTS; PR00011; GLAMININ G_2; PR00018; EGF; 4.

SMART; SM00189; GLA; 3.

SMART; SM00089; GLA; 2.

SMART; SM00089; GLA; 2.

SMART; SM00209; EGF_1; UNIO

PROSITE; PS00010; EGF_2; 3.

PROSITE; PS01016; EGF_2; 3.

PROSITE; PS01016; EGF_2; 3.

PROSITE; PS01016; EGF_3; 4.

PROSITE; PS00117; GLA_1; UNIM

PROSITE; PS00117; GLA_2; 1.

PROSITE; PS0099; GLA_2; 1.

PROSITE; PS0099; GLA_3; 1.

PROSITE; PS50998; GLA_3; 0.

Calcium; EGF_1ke domain; Sisignal
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Stenflo J., Wydro R.;
"Isolation and sequence of the cDNA for human pro
of blood coagulation.";
Proc. Natl. Acad. Sci. U.S.A. 83:6716-6720(1986).
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NUCLEOTIDE SEQUENCE.
MEDLINE=86313649; PubMed=2944113;
Nackowski W., Cohen E.,
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Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Copyrighted by the UniProt Consortium, Distributed under the Creative Commons
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PF00054;
PF02210;
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rPro; IPR000294;
; PF00008; EGF; 2
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        3; PS00010; ASX HYDROXYL; 3.
3; PS01186; EGF 1; UNKNOWN 1.
3; PS01186; EGF 2; 3.
3; PS50026; EGF 3; 4.
3; PS001187; EGF CA; 2.
3; PS01187; EGF CA; 2.
3; PS00011; GLA 1; UNKNOWN 1.
3; PS50926; LAM G DOMAIN; 2.
1; EGF - like domain; Signal.
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                                                                                                                                                                                                                                                                                                      ISDO HUMAN PRELIMINARY;

Q9NSDO_HUMAN PRELIMINARY;

Q9NSDO;

O1-OCT-2000, integrated into

O1-CT-2000, sequence versio

O7-FEB-2006, entry version 1
                                           TISSUE=Liver;
Wydro R., Coh
Dahlback B.;
                                                                                                                                                         NCBI_TaxID=9606;
                                                                                                                                                                                                            Mammalia;
                                                                                                                                                                                                                                  Homo sapiens (Human
Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CHAIN
NON_TER
                        Submitted (FEB-1992)
                                                                                                                       NUCLEOTIDE
                                                                                                                                                                                                                                                                                Protein S precursor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PVLTFAGGLPDVPVTSAPVTAFYRGCMTLEVNRRLLDLDEAAYKHSDITAHSCPPV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GTESTWEVEVVAHIRPAADTGVLFALWAPDLRAVPLSVALVDYHSTKKLKKQLVVLAVEH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RLDGCMRSWNWLNGEDTTIQETVKVNTRMQCFSVTERGSFYPGSGFAFYSLDYMRTPLDV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MVSVEELEHSISIKIAKEAVMDINKPGPLFKPENGLLETKVYFAGFPRKVESELIKPINP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AEFDFRTFDPEGILLFAGGHQDSTWIVLALRAGRLELQLRYNGVGRVTSSGPVINHGMWQ 380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A-DSEACGEARCKNLPGSYSCLCDEGFAYSSQEKACRDVDECLQGRCEQVCVNSPGSYTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TAL-ALMEIKVC-DGQEHVVTVSLRDGEATLEVDGTRGQSEVSAAQLQERLAVLERHLRS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SSAEGWHVNVTLNIRPSTGTGVMLALVSGN-NTVPFAVSLVD--STSE-KSODILLSVEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REDGCIRSWNLMKQGASGIKEIIQEKQNKHCLVTVEKGSYYPGSGIAQFHIDYN----NV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AEFDFRTYDSEGVILYAESIYHSAWLLIALRGGKIEVQLKNEHTSKITTGGDVINNGLWN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    YCDGKKGFKLAQDQKSCEVVSVCLPLNLDTKYELLYLAEQFAGV-VLYLKFRLPEISRFS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HCDGRGGLKLSQDMDTCEDILPCVPFSVAKSVKSLYLGRMFSGTPVIRLRFKRLQPTRLV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LVCLRSFQTGLFTAARQSTNAYPDLRSCVNAIPDQCSPLPCNEDGYMSCKDGKASFTCTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LDCINKY-----GSPYTKNS--GFATCVQNLPDQCTPNPCDRKGTQACQDLMGNFFCLC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LPAREATQFLRPRQRRAFQVFEEAKQGHLERECVEELCSREEAREVFENDPETDYFYPRY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SLKPSICGTAVCKNILGDFECECPEGYRYNLKSKSCEDIDECSENMCAQLCVNYPGGHTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KPGWQGEKCEFDINECKDPSNINGGCSQICDNTPGSYHCSCKNGFVMLSNKKDCKDVDEC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         KAGWGGRLCDKDVNEC---SQENGGCLQICHNKPGSFHCSCHSGFELSSDGRTCQDIDEC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LSKQQASQVL-VRKRRANSLLEETKQGNLERECIEELCNKEEAREVFENDPETDYFYPKY
                                                                 Cohen E.,
                                                                                                                                                                                                               Eutheria;
                                                                                                                  SEQUENCE
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                                                                                                                                                                                                                                                                                                      integrated into UniProtKB/TrEMBL
sequence version 1.
entry version 19.
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                                                                                                                                                                                                                                                             (Human)
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                                                                    Dackowski W.,
                                                                                                                                                                                                          Euarchontoglires; Primates; Catarrhini;
                                                                                                                                                                                                                                    Chordata;
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Pred. No. 2.2e-
27; Mismatches
                     EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                  Craniata; Vertebrata; Euteleostomi
                                                                                                                                                                                                                                                                                                                                                                                                   PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LEFRVNRNNLELSTPLKIETISHEDLQRQLAVLDKAMKA
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                                                                    Stenflo
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2.2e-99;
                                                                 J.
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                        databases
                                                                    Lundwall
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                                                                                                                                                                                                               Hominidae;
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SOUTH TAKE
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Best Local S
Matches 290
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Pfam; PF000084; EGF CA; 1.

Pfam; PF00054; Laminin G_1; 1

Pfam; PF00054; Laminin G_2; 1

Pfam; PF00210; Laminin G_2; 1

Pfam; PF002210; Laminin G_2; 1

SMART; SM00189; EGF CA; 3.

SMART; SM00189; GLA; 1.

SMART; SM00282; Laminin G_2; 2.

PROSITE; PS00010; ASX HYDROXY.

PROSITE; PS00022; EGF 1; UNKNUPROSITE; PS01186; EGF 2; 3.

PROSITE; PS01186; EGF 3; 4.

PROSITE; PS01187; EGF CA; 2.

PROSITE; PS00011; GLA 1; UNKNUPROSITE; PS00011; GLA 2; 1.

PROSITE; PS00098; GLA 2; 1.

PROSITE; PS00098; GLA 2; 1.

PROSITE; PS00029; LAM_G_DOMAI.

SIGNAL 1 15

CHAIN 16 650

SEQUENCE 650 AA; 72480 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 InterPro; IPR013320; C
InterPro; IPR006210; E
InterPro; IPR000742; E
InterPro; IPR001881; E
InterPro; IPR013091; E
InterPro; IPR013030; E
InterPro; IPR012303; E
InterPro; IPR002383; G
InterPro; IPR002383; G
InterPro; IPR0126791; I
InterPro; IPR012680; L
InterPro; IPR012680; L
InterPro; IPR010294; V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; X12892; CHSSP; P00740; 1
SMR; Q9NSD0; 17
Ensembl; ENSG00
GO; GO:0005576;
GO; GO:0005509;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Distributed under the Creative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Copyrighted by the UniProt Consortium,
                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GO:0005509;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      pmbl; ENSG00000184500; Homo sapiens.
GO:0005576; C:extracellular region;
                                                                                                                                 204
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 302
                          324
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                                                                                                       183
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                                                                                                                                                                                                                                                                  4.
                                                                                                                                                                                                                                                                                                                                                                                                                       3; PS00010; ASX HYDROXYL; 3.
5; PS00022; EGF 1; UNKNOWN 1.
5; PS01186; EGF 2; 3.
6; PS50187; EGF CA; 2.
7; PS001187; EGF CA; 2.
7; PS00011; GLA 2; 1.
7; PS50025; LAM_G_DOMAIN; 2.
                                                                                                                                                                                                                                                                               GRGGLKLSQDMDTCCDILPCVPFSVAKSVKSLYLGRMFSGTPVIRLRFKRLQPTRLVAEF
                                                                                                                       SEACGEARCKNILPGSYSCLCDEGFAYSSQEKACRDVDECLQGRCEQVCVNSPGSYTCHCD
                                                                                                                                                           WQGEKCEFDINECKDPSNINGGCSQICDNTPGSYHCSCKNGFVMLSNKKDCKDVDECSLK
                                                                                                                                                                                  WGGRLCDKDVNEC---
                                                                                                                                                                                                               LRSFQTGLFTAARQSTNAYPDLRSCVNAIPDQCSPLPCNEDGYMSCKDGKASFTCTCKPG
                                                                                                                                                                                                                                       INKY-----GSPYTKNS--GFATCVQNLPDQCTPNPCDRKGTQACQDLMGNFFCLCKAG
DFRTYDSEGVILYAESIYHSAWLLIALRGGKIEVQLKNEHTSKITTGGDVINNGLWNMVS
                        DFRTFDPEGILLFAGGHQDSTWIVLALRAGRLELQLRYNGVGRVTSSGPVINHGMWQTIS
                                                                                                        PSICGTAVCKNILGDFECECPEGYRYNLKSKSCEDIDECSENMCAQLCVNYPGGHTCYCD
                                                    GKKGFKLAODOKSCEVVSVCLPLNLDTKYELLYLAEOFAGV
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650
                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                         15
650
72480 <sub>1</sub>
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EGF Ca bd.

EGF Ca bd 2.

EGF like reg.

EGF like reg.

EGF like reg.

Laminin G.

Laminin G.

Laminin G.

Laminin G.

Laminin G.

Vitk dep GLA.
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                                                                                                                                                                                -SQENGGCLQICHNKPGSFHCSCHSGFELSSDGRTCQDIDECA-D
                                                                                                                                                                                                                                                                                                                    127;
                                                                                                                                                                                                                                                                                                                                                                         MW;
                                                                                                                                                                                                                                                                                                                                                                                      Potential. Potential.
                                                                                                                                                                                                                                                                                                                 Pred. No. 3.1e
7; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mRNA.
                                                                                                                                                                                                                                                                                                                                  Score 1514;
Pred. No. 3
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PROS_I
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07-WAR-2006, entry version 70.
Vitamin K-dependent protein S precursor.
          EMBL; M13044; AAA30757.1;
EMBL; X12891; CAA31382.1;
PIR; A24759; KXBOS.
HSSP; P00740; ICFH.
                                                                                                                       -:- SUBCELLULAR LOCATION: Secreted protein.
-!- TISSUE SPECIFICITY: Plasma.
-!- SIMILARITY: Contains 4 EGF-like domains.
-!- SIMILARITY: Contains 1 Gla (gamma-carboxy-glutamate)
-!- SIMILARITY: Contains 2 laminin G-like domains.
                                                                                                                                                                                                                                 Dahlback B., Lundwall A., Stenflo J., "Localization of thrombin cleavage si of bovine protein S.", J. Biol. Chem. 261:5111-5115(1986) -i- FUNCTION: Anticoagulant plasma pr
                                                                                                                                                                                                                                                                                                                                                                                                                                            NUCLEOTIDE SEQUENCE [MRNA].

MEDLLINE=8623400; PubMed=2940598;

Mahlback B., Lundwall A., Stenflo J.;

Pahlback B., Lundwall A., Stenflo J.;

Primary structure of bovine vitamin K-dependent Proc. Natl. Acad. Sci. U.S.A. 83:4199-4203(1986).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Crania
Mammalia; Eutheria; Laurasiatheria;
Pecora; Bovidae; Bovinae; Bos.
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01-APR-1988,
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Dahlback B.;
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                                                                                                                                                                                                      FUNCTION: Anticoagulant plasma protein; it is a cofactor to activated protein C in the degradation of coagulation factors and VIIIa. It helps to prevent coagulation and stimulating
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Pfam; PF00008; EGF; 2.1.

R Pfam; PF007645; EGF; CA; 3.

R Pfam; PF00594; Laminin G 1; 1.

R Pfam; PF00594; Laminin G 2; 2.

R Pfam; PF00054; Laminin G 2; 2.

R Pfam; PF00001; GLABLOOD.

R SMART; SM00181; EGF; 4.

SMART; SM00179; EGF; CA; 3.

SMART; SM00179; EGF; CA; 3.

SMART; SM00179; EGF; CA; 1.

R SMART; SM00189; GLA; 1.

PROSITE; PS00010; ASX HYDROXYL; 4.

PROSITE; PS01186; EGF 2; 3.

PROSITE; PS01186; EGF 3; 3.

PROSITE; PS01187; EGF-CA; 3.

PROSITE; PS00011; GLA 1; 1.

R PROSITE; PS00011; GLA 1; 1.

PROSITE; PS00025; LAMG GDOMAIN; 2.

PROSITE; PS00025; LAMG; Direct protein set Blood coagulation; Repeat; Signal; Zymogen.

FYGINAL 25

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N-linked (GlcNA)
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Vitamin K-dependent p
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3675
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

| 23 | 22 | 21 | 20 | 19 | 18 | 17 | 16 | 15 | 14 | 13 | 12 | 11 | 10 | 9 | 8 | 7 | 0 | л | 4. | ω | 2 | 1 | Result |
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| ADE41298 | ADD37090 | ADD11297 | ADD10337 | ABB95446 | ABB84840 | AAY57382 | AAB33459 | AAY29793 | AAW46462 | ADQ67207 | ADD48757 | AEE87023 | ADV91450 | ADT91779 | ADN60275 | ADM40825 | ADL83231 | AAY57383 | AAY29794 | AAW46463 | AAR99414 | ADM40823 | Ð |
| Human | Human | | | | Abb84840 Human PRO | Amino | | Aay29793 Murine gr | Murine | Novel | Add48757 Human Pro | Aee87023 Human che | Human | | Human | Human | Human | Amino | Human | Human | Aar99414 Human gas | Adm40823 Human Gas | |

| Human gro | Adv91465 | ADV91465 | 9 | 227 | 35.3 | 1297 | 45 |
|-----------|-----------|----------|----|-----|------|--------|-----|
| Human pro | Aar70728 | AAR70728 | N | 675 | 39.1 | 1437.5 | 44 |
| Rat Prote | Add48898 | ADD48898 | 7 | 675 | 39.4 | 1448 | 43 |
| Bovine Pr | Aap81136 | AAP81136 | ш | 675 | 40.2 | 1479 | 42 |
| Vitamin K | Aar31875 | AAR31875 | N | 635 | 41.1 | 1510.5 | 41 |
| Mature hu | Aar72350 | AAR72350 | N | 635 | 41.1 | 1511.5 | 40 |
| Human rec | Aap70083 | AAP70083 | _ | 676 | 41.2 | 1515.5 | 9 |
| Human Pro | Add48900 | ADD48900 | 7 | 650 | 41.3 | 1516 | 38 |
| Human pro | Aap81137 | AAP81137 | μ. | 650 | 41.3 | 1516 | 37 |
| Human sof | Adq17650 | ADQ17650 | æ | 650 | | 1532 | 36 |
| Human Pro | Ade62065 | ADE62065 | 7 | 676 | | 1543 | 35 |
| Human Pro | Add46140 | ADD46140 | 7 | 676 | 42.0 | 1543 | 34 |
| Amino aci | Aay57384 | AAY57384 | ω | 676 | 42.0 | 1543 | 33 |
| Human pro | Aay29795 | AAY29795 | N | 676 | 42.0 | 1543 | 32 |
| Human pro | Aaw46464 | AAW46464 | N | 676 | 42.0 | 1543 | 31 |
| Human pro | Adz14050 | ADZ14050 | ø | 676 | 42.1 | 1547.5 | 30 |
| Human GAS | Adt 91805 | ADT91805 | æ | 676 | 42.1 | 1547.5 | 29 |
| PRO poly | Adp24054 | ADP24054 | œ | 676 | 42.1 | 1547.5 | 28 |
| Human can | Abo84421 | ABO84421 | œ | 518 | 71.3 | 2622 | 27 |
| Mouse can | Abo84420 | ABO84420 | 8 | 703 | | 2931 | 26 |
| Human PRC | Adk82826 | ADK82826 | 8 | 673 | 80.4 | 2955.5 | 25 |
| Human PKC | AGD43481 | ADH43481 | a | 0/0 | 000 | 6,000 | 7.4 |

ALIGNMENTS

RESULT 1

growth arrest specific gene 6; Gas6; epitope tag; variant Gas6; anti-apoptotic; Gas6 inhibitor; Gas6-dependent receptor phosphorylation; receptor internalisation; cell proliferation; cell apoptosis prevention; signaling molecule; cell marker; human. 08-APR-2004. WO2004029209-A2 Human Gas6 protein with a C-terminal epitope tag SEQ ID NO:1. 01-JUL-2004 Homo sapiens ADM40823; ADM40823 standard; (first entry) protein; 989 ₿

(CENZ) CENTOCOR INC.

24-SEP-2002; 2002US-0413157P 24-SEP-2003; 2003WO-US030330.

Yang J, Chen C, 표 B Andrade-Gordon P, Jordan

WPI; 2004-316097/29.

New epitope-tagged Growth Arrest Specific Gene 6 (Gas6) proteins for manufacturing medicines or for identifying inhibitors of Gas6 and its

Claim 2; SEQ ID NO 1; 28pp; English

The present invention describes an isolated growth arrest specific gene 6 (Gas6) polypeptide, or its variant, with an epitope tag fused at the C-terminus. The epitope tag comprises a Flag sequence of 8 amino acids (SEQ ID NO:2, ADM40824), or a polyhistidine sequence. The variant Gas6 polypeptide has 75% identity to native human Gas6 protein. Also described: (1) a recombinant DNA molecule encoding the amino acid sequence of the polypeptide described above; (2) a vector comprising the above nucleic acid; (3) a host cell comprising the vector, where the host cell is of mammalian origin; and (4) a composition comprising the

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RESULT 2
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DT 04-I
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Pred. No. 7.1e-248;
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Axl receptor ligand; human protein S homologue; myeloid cell proliferation; myeloid malignancy;
                                       recombinant production
sapiens.
                                                           growth factor;
                                                                                  factor;
                                                             prevention;
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23-JUL-1996

94US-00282141

29-JUL-1994; 94US-00282141

(AMGE-) AMGEN INC. SCHNEIDER G

Manfioletti G, Varnum BC, Schneider ņ, Avanzi g

WPI; 1996-353825/35 N-PSDB; AAT41544.

DNA encoding gas6, the axl receptor myeloid cells and malignancies. ligand 1 useful 6 regulate growth

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Col 19-22; 36pp; English.

AAR99414 is the gas6 protein (from the growth arrest specific gene number 6). The gas6 protein has homology to human protein S which functions as a cofactor in a protease cascade that regulates coagulation. Gas6 expression, as with gas1 and gas2, is associated with cell growth arrest which suggests a possible role of gas6 in the regulation of cell growth. In fact the gas6 protein is an ax1 receptor (ax1r) ligand and a growth factor for any cells expressing ax1r e.g. bone marrow, spleen, thymus, ovary, heart, intestine and lung cells. The ax1 receptor is involved in myeloid cell proliferation and myeloid malignancies, so gas6 may regulate growth of such cells

Sequence 678 AA;

Conservative

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98.7%; 100.0%;

Score 3628; DB 2; Pred. No. 1.3e-244; Mismatches

Length Indels

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RESULT 3
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       Medium for culturing human Schwann cells receptor activator and other mitogens.
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                                    WPI; 1998-129864/12.
                                                                                        10-MAY-1995;
                                                                                                          10-MAY-1995;
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Disclosure; Fig 2; 51pp; English

The present sequence represents a human growth arrest specific-gene 6 (gas6) protein which is able to activate the Rea and Axl receptor. Ree is a receptor tyrosine kinase that is preferentially expressed in the adult brain. Gas6 is a vitamin K dependent protein which may play a role in the regulation of a protease cascade relevent in growth regulation. The protein is used in a serum-free culture medium for culturing human schwann cells, which does not support fibroblast growth. Gas6 functions as a mitogenic agent which is a Res/Axl receptor activator. A second mitogenic agent e.g. insulin or an erbB receptor activator such as heregulin, is also required to raise cAMP levels and enhance survival or proliferation of human Schwann cells. The culture medium additionally comprises a molecule or composition that provides Fe ions to the Schwann cells, vitamin E, a protease inhibitor, and progesterone. The Schwann cells can be used to treat patients with nervous system injuries

Sequence 678 AA;

Similarity

98.7%; So lilarity 100.0%; I Conservative 0;

Score 3628; DB 2; 1 Pred. No. 1.3e-244; 0; Mismatches 0;

Length Indels

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rse receptor protein tyrosine kinase; Rse ligand; Rse-L; gas6; growth arrest-specific gene 6; proliferation; differentiation; glial cell; Schwann cell; fusion protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                A method has been developed for activating the Rse receptor on a cell. The method comprises exposing the receptor to the exogenous growth arrei-specific gene 6 (gas6) polypeptide. The method is useful for enhancing cell proliferation and cell differentiation. The present sequence
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The invention relates to a method for treating nervous system injuries in mammals by administering human Schwann cells (SC) that have been cultured in serum-free medium. The serum-free medium is a nutrient solution supplemented with two mitogens, one of which, is a Res/Axl receptor activator, to increase survival and proliferation of SC. The method is used to treat (or prevent) central or peripheral nervous system injury, e.g. traumatic, ischemic, malignant, infectious or degenerative lesions, or lesions associated with nutritional deficiencies, systemic disease, toxins or demyelinization. Culturing cells in the specified medium allows proliferation of adult SC for use in autologous transplants. The present sequence represents a human gas6 (h gas6) protein, having 44% sequence identity to human protein S. Gas6 is demonstrated to be a potent growth/survival factor for SCs in defined serum-free culture

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Disclosure; Fig 2; 52pp; English.

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 WO2004024097-A2
                         Homo sapiens
                                                  immune-mediated inflammatory
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LERECVEELCSREEAREVFENDPETDYFYPRYLDCINKYGSPYTKNSGFATCVQNLPDQC

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LTVGGI PFHEKDLVQPINPRLDGCMRSWNWLNGEDTTIQETVKVNTRMQCFSVTERGSFY

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YNGVGRVTSSGPVINHGMWQTISVEELARNLVIKVNRDAVMKIAVAGDLFQPERGLYHLN FSGTPVIRLRFKRLQPTRLVAEFDFRTFDPEGILLFAGGHQDSTWIVLALRAGRLELQLR FSGTPVIRLRFKRLQPTRLVAEFDFRTFDPEGILLFAGGHQDSTWIVLALRAGRLELQLR ECLQGRCEQVCVNSPGSYTCHCDGRGGLKLSQDMDTCEDILPCVPFSVAKSVKSLYLGRM ECLQGRCEQVCVNSPGSYTCHCDGRGGLKLSQDMDTCEDILPCVPFSVAKSVKSLYLGRM SCHSGFELSSDGRTCQDIDECADSEACGEARCKNLPGSYSCLCDEGFAYSSQEKACRDVD SCHSGFELSSDGRTCQDIDECADSEACGEARCKNLPGSYSCLCDEGFAYSSQEKACRDVD TPNPCDRKGTQACQDLMGNFFCLCKAGWGGRLCDKDVNECSQENGGCLQICHNKPGSFHC TPNPCDRKGTQACQDLMGNFFCLCKAGWGGRLCDKDVNECSQENGGCLQICHNKPGSFHC

YNGVGRVTSSGPVINHGMWQTISVEELARNLVIKVNRDAVMKIAVAGDLFQPERGLYHLN

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301 301

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The present invention relates to PRO proteins and their coding sequences.

CC The PRO proteins are useful for diagnosing and treating a B cell related disorder, e.g. X-linked infantile hypogammaglobulinemia, polysaccharide antigen unresponsiveness, selective IgA deficiency, selective IgM deficiency, selective deficiency of IgG subclasses, immunodeficiency with Chyper IgM, transient hypogammaglobulinemia of infancy, Burkitt's CC hypersensitivity, rheumatoid arthritis, autoimmune mediated haemolytic anaemia, myasthenia gravis, hypoadarencocriticism, glomerulonephritis, or CC anaemia, myasthenia gravis, hypoadarencocriticism, glomerulonephritis, or CC antylosing spondylitis. The PRO proteins are also useful for preparing a condition that is responsive to the PRO coding sequences are useful as hybridization probes in chromosome and CC gene mapping, in preparing PRO proteins, or in generating transgenic animals or knockout animals, which in furn are useful in the development
      Best Local Similarity Matches 678; Conserv
                      Query Match
Best Local (
                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                New PRO polypeptide, useful for diagnosing and treating disorder, e.g. Burkitt's lymphoma, rheumatoid arthritis, mediated hemolytic anemia, myasthenia gravis or ankylosi
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                 The present invention describes an isolated growth arrest specific gene 6 (Gas6) polypeptide, or its variant, with an epitope tag fused at the C-terminus. The epitope tag comprises a Flag sequence of 8 amino acids (SEQ ID NO:2, ADM40824), or a polyhistidine sequence. The variant Gas6 polypeptide has 75% identity to native human Gas6 protein. Also described: (1) a recombinant DNA molecule encoding the amino acid sequence of the polypeptide described above; (2) a vector comprising the sequence of the polypeptide described above; (2) a vector, where the host cell is of mammalian origin; and (4) a composition comprising the polypeptide and a pharmaceutical carrier. Gas6 has anti-apoptotic activity. The composition is useful for manufacturing medicines or for identifying inhibitors of Gas6 and its receptors. The Gas6 polypeptide may also be used for Gas6-dependent receptor phosphorylation, receptor infernalisation, cell proliferation, prevention of cell apoptosis, or induction of signaling molecules or cell markers. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               growth arrest specific gene 6; Gas6; epitope tag; variant Gas6; anti-apoptotic; Gas6 inhibitor; Gas6-dependent receptor phosphorylation; receptor internalisation; cell proliferation; cell apoptosis prevention; signaling molecule; cell marker; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New epitope-tagged Growth Arrest Specific Gene 6 (Gas6) proteins manufacturing medicines or for identifying inhibitors of Gas6 and
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                                                                                                                                                                                                                                                                                                                                               The present invention describes a method for identifying a compound that modulates angiogenesis or tumourigenesis. The method comprises: (a) contacting the compound with angiogenesis polypeptide, e.g. Ax1, tubulin cofactor D, transglutaminase 2, cytosine deaminase, peptidase M41 (paraplegin), CD13 aminopeptidase, PPK-1, zip kinase, Gas6, SRm160, non-muscle myosin heavy chain, calmodulin 2, novel symporter, novel semaphorin, novel zinc finger helicase (FLJ22611), plexin-A2, deoxycytidylate deaminase or novel sugar transporter; (b) determining the functional effect of the compound upon the angiogenesis polypeptide or the physical effect of the compound upon the target polypeptide or its fragment or inactive variant; and (c) determining the chemical or polypeptide or its fragment or inactive variant, thus identifying a compound that modulates cell cycle arrest. Also described is a method of modulating angiogenesis in a subject. The angiogenesis or tumourigenesis modulating compound has antiangiogenic, cytostatic, cerebroprotective, vasotropic, antisheritlity and cardiant activities, and can be used in antibody, antisense and RNA interference (RNAi) therapies. The methods angiogenesis, cancer, stroke, infertility and heart disease. The present sequence represents a human growth arrest-specific 6 (GAS6) protein, which is used in the exemplification of the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Identifying a compound that modulates angiogenesis or tumorigenesis, useful in diagnosing and treating angiogenesis, cancer, stroke, infertility and heart disease, comprises contacting the compound with angiogenesis polypeptide.
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                                                                                                                Identifying a compound for treating osteoarthritis (OA) comprises contacting a test compound to a reaction mixture that comprises a polypeptide member of the TYRO3 subfamily of receptor tyrosine kiand a ligand to the polypeptide.
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Pred. No. 1.3e-244;
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                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                     (Gas6) polypeptide. The invention also relates to a recombinant DNA molecule encoding the polypeptide, an antibody or antigen binding fragment capable of binding to the polypeptide and a labeled antibody generated against Gas6. The Gas6 polypeptide is useful for generating antibodies against Gas6, which involves immunization with the polypeptide or screening recombinant antibodies with the polypeptide. The antibodies are useful for detecting Gas6 polypeptide in a sample suspected of containing Gas6. Compositions containing the polypeptide are useful for preventing or treating cardiovascular diseases resulting from a dysfunction in a mammal. The compositions are also useful for treating cancer, atherosclerosis, sepsis, glomerulosclerosis, diabetes, rheumatoid arthritis, HIV infection, osteoarthritis, osteoporosis and multiple sclerosis. This sequence represents the human Gas6 polypeptide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            multiple sclerosis; cardiovascular-gen.; antirheumatic; antiarthritic; anti-HIV; cytostatic; neuroprotective; thrombolytic; cardiant; cerebroprotective; vasotropic; antiarteriosclerotic; antibacterial; immunosuppressive; nephrotropic; antidiabetic; osteopathic;
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N-PSDB; AEE86958.
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Investigating response to anti-cancer chemotherapy comprises determining the expression level of gene(s) of interest in a biopsy sample.

IJ NO 107; 248pp; English

This invention relates to a novel method of investigating response to anti-cancer chemotherapy. The method comprises determining, in a biopsy sample taken from a neoplastic lesion after the onset of a chemotherapy schedule, the expression level of up to 65 genes included in the specification. The invention may be useful for the development of compounds with a cytostatic activity whilst the disclosed sequences may prove useful for gene therapy. The method is useful for investigating response to anti-cancer chemotherapy. The method, composition, and kit are useful or predicting, diagnosing, prognosing, preventing, and treating malignant neoplasia and breast cancer. The present sequence is that of the protein encoded by a human gene related to the response to cancer chemotherapy. The gene was claimed for use in the method of the

Sequence 678 AA;

Match

Similarity

Length

678;

Local Sim: hes 678; 541 481 481 421 421 361 361 301 301 241 241 181 121 181 121 61 61 KHSDITAHSCPPVEPAAA AAQLQERLAVLERHLRSPVLTFAGGLPDVPVTSAFVTAFYRGCMTLEVWRRLLDLDEAAY DYHSTKKLKKQLVVLAVEHTALALMEI KVCDGQEHVVTVSLRDGEATLEVDGTRGQSEVS PGSGFAFYSLDYMRTPLDVGTESTWEVEVVAHIRPAADTGVLFALWAPDLRAVPLSVALV LTVGGIPFHEKDLVQPINPRLDGCMRSWNWLNGEDTTIQETVKVNTRMQCFSVTERGSFY LTVGGIPFHEKDLVQPINPRLDGCMRSWNWLNGEDTTIQETVKVNTRMQCFSVTERGSFY FSGTPVIRLRFKRLQPTRLVAEFDFRTFDPEGILLFAGGHQDSTWIVLALRAGRLELQLR ECLQGRCEQVCVNSPGSYTCHCDGRGGLKLSQDMDTCEDILPCVPFSVAKSVKSLYLGRM SCHSGFELSSDGRTCQDIDECADSEACGEARCKNLPGSYSCLCDEGFAYSSQEKACRDVD TPNPCDRKGTQACQDLMGNFFCLCKAGWGGRLCDKDVNECSQENGGCLQICHNKPGSFHC LERECVEELCSREEAREVFENDPETDYFYPRYLDCINKYGSPYTKNSGFATCVQNLPDQC MAPSLSPGPAALRRAPQLLLLLLAAECALAALLPAREATQFLRPRQRRAFQVFEEAKQGH KHSDITAHSCPPVEPAAA DYHSTKKLKKQLVVLAVEHTALALMEIKVCDGQEHVVTVSLRDGEATLEVDGTRGQSEVS PGSGFAFYSLDYMRTPLDVGTESTWEVEVVAHIRPAADTGVLFALWAPDLRAVPLSVALV **FSGTPVIRLRFKRLOPTRLVAEFDFRTFDPEGILLFAGGHQDSTWIVLALRAGRLELQLR** MAPSLSPGPAALRRAPQLLLLLLAAECALAALLPAREATQFLRPRQRRAFQVFEEAKQGH ECLOGRCEQVCVNSPGSYTCHCDGRGGLKLSQDMDTCEDILPCVPFSVAKSVKSLYLGRM SCHSGFELSSDGRTCQDIDECADSEACGEARCKNLPGSYSCLCDEGFAYSSQEKACRDVD TPNPCDRKGTQACQDLMGNFFCLCKAGWGGRLCDKDVNECSQENGGCLQICHNKPGSFHC LERECVEELCSREEAREVFENDPETDYFYPRYLDCINKYGSPYTKNSGFATCVQNLPDQC 98.7%; Score 3628; DB 10; ilarity 100.0%; Pred. No. 1.3e-244; Conservative 0; Mismatches 0; 678 Indels 0 Gaps 660 600 600 540 480 480 420 420 360 360 300 300 240 240 180 180 120 120 60

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                                                                                                      CC which is differentially regulated in an animal subjected to pain and a CC kit to perform the method, an array, a method for identifying an agent that increases or decreases the expression of the polynucleotide sequence CC that is differentially expressed in neuronal tissue of a first animal CC subjected to pain, a method for identifying a compound which regulates the expressed in an animal subjected to pain, a method for identifying a CC compound that regulates the activity of one or more of the polynucleotides, a method for producing a pharmaceutical composition, a CC method for identifying a compound or small molecule that regulates the CC extivity in an animal of one or more of the polynucleotide of the specification, a method for identifying a compound useful in treating pain and a pharmaceutical composition comprising the one or more CC modulates its activity is useful for preparing a medicament for treating CC pain and a pharmaceutical nerve injury (Chung), chronic constriction conjury (CCI) and spared nerve injury (SNI)) in an animal (e.g. gene CC the specification) which is differentially expressed during pain. CC Note: The sequence data for this patent did not form part of the printed Specification, but was obtained in electronic form directly from WIPO at CC ftb.wipo.inf/mb/rubh/libled nor semenars.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention discloses a composition comprising two or more isolated rat or human polynucleotides or a polynucleotide which represents a fragment, derivative or allelic variation of the nucleic acid sequence. Also claimed are a vector comprising the novel polynucleotide, a host cell comprising the vector, a method for identifying a nucleotide sequence which is differentially regulated in an animal subjected to pain and a which is differentially regulated in an animal subjected to pain and a possible of the composition o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New composition comprising two or more isolated polypeptides, useful preparing a medicament for treating pain in an animal.
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01-NOV-2001; 2001US-0346382P
26-NOV-2001; 2001US-0333347P
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Query Match

Score 3628;

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Length 679;

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                                                                                                                                                                                                      osteopathic; neuroprotective; nootropic; antiparkinsonian; cytostatic; gene therapy; diagnostic marker; morbid state; osteoporosis; neurological disease; Alzheimer's disease; Parkinson's disease; dementia;
                                                                                                                                                                                                                                                                                                              Novel human
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Best Local S
Matches 584
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    are useful as diagnostic markers or therapeutic target for the diseases or morbid states. They are also useful for treating osteoporosis, neurological diseases, Alzheimer's diseases, Parkinson's diseases, dementia and various cancers. This sequence corresponds to a protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to 2495 novel polynucleotides (I) and their encoded polypeptides, sequences hybridizing to these nucleotides, sequences encoding partial polypeptides and sequences having 70% or 90% identity to the nucleotide and protein sequences. The nucleotides and polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Isogai T,
Yamamoto :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
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                                                                                                                                                                                                                            KVNRDAVMKIAVAGDLFQPERGLYHLNLTVGGIPFHEKDLVQPINFRLDGCMRSWNWLNG
                                                                                                                                                                                                                                             KVNRDĀVMKIAVAGDLFQPERGLYHLNLTVGGIPFHEKDLVQPINPRLDGCMRSWNWLNG
                                                                                                                                                                                                                                                                                                        LLFAGGHQDSTWIVLALRAGRLELQLRYNGVGRVTSSGPVINHGMWQTISVEELARNLVI
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          APVTAFYRGCMTLEVNRRLLDLDEAAYKHSDITAHSCPPVEPAAA
                                                                                                           RPAADTGVLFALWAPDLRAVPLSVALVDYHSTKKLKKQLVVLAVEHTALALMEIKVCDGQ
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Isono Y,
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Pred. No. 5.5e-211;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence represents a murine growth arrest specific-gene 6 (gas6) protein which is able to activate the Rsa and Axl receptor. Rse is a receptor tyrosine kinase that is preferentially expressed in the adult brain. Gas6 is a vitamin K dependent protein which may play a role in the regulation of a protease cascade relevent in growth regulation. The protein is used in a serum-free culture medium for culturing human schwann cells, which does not support fibroblast growth. Gas6 functions as a mitogenic agent which is a Rse/Axl receptor activator. A second mitogenic agent e.g. insulin or an erbB receptor activator such as heregulin, is also required to raise CAMP levels and enhance survival or proliferation of human Schwann cells. The culture medium additionally comprises a molecule or composition that provides Fe ions to the Schwann cells, vitamin E, a protease inhibitor, and progesterome. The Schwann cells can be used to treat patients with nervous system injuries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 673
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Growth arrest specific-gene 6; gas6; Rsa; Axl; receptor; mitogenic agent receptor tyrosine kinase; regulation; protease cascade; insulin; growth regulation; serum-free culture medium; human; Schwann cell; receptor activator; exbB receptor; heregulin; cAMP level; proliferation;
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PGPAA-ALGTALLLLLASESSHTVLLRAREAAQFLRPRQRRAYQVFEBAKQGHLERECV
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2: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US08_PUBCOMB.pep:*

3: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US09_PUBCOMB.pep:*

4: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10A_PUBCOMB.pep:*

5: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*
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summaries
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Sequence 3, Appli
Sequence 41949, Ap
Sequence 48, Appl
Sequence 31, Appl
Sequence 31, Appl
Sequence 30, Appl
Sequence 37, Appl
Sequence 66, Appli
Sequence 67, Appli
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361 YNGVGRVTSSGPVINHGMWQTISVEELARNLVIKVNRDAVMKIAVAGDLFQPERGLYHLN 420
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| 361 YNGV | 301 FSGT 301 FSGT | 241 BCLC 241 ECLC | 181 SCHS 181 SCHS | 121 TPNPC | 61 LERE 61 LERE | 1 MAPS 1 MAPS | atch cal Simi 686; | SULT 1 Sequence 1, Application 1 Publication No. US2004011 GENERAL INFORMATION: APPLICANT: JOHNSON & JO! TITLE OF INVENTION: EPI' TITLE OF INVENTION: EPI' TITLE OF INVENTION: PRICE CURRENT APPLICATION NUM CURRENT FILING DATE: 2012 PRIOR APPLICATION NUMBER PRIOR FILING DATE: 2002- NUMBER OF SEQ ID NOS: 3 SOFTWARE: Patentin versi SEQ ID NO 1 LENGTH: 686 TYPE: PRT ORGANISM: Homo sapiens -10-671-054-1 | | 35.5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 |
| YNGVGRVTSS | PVIRLR PVIRLR | ,QGRCEQV ,QGRCEQV | GFELS | CDRKGT | LERECVEELC | SLSPGPA | larity Conserv | lication US200401 TION: EDITION: EPITION: EPITION: EPITION: EPITION: PRITION: PRITION: PRITION: PRITION: PRITION: PRITION NUMBE: 2002 ICN NUMBE: 2002 ICN VORBE: 2002 ICN VORBE: 2002 STILL VORB | | ,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,, |
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RESULT 2
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TITLE OF INVENTION: EPITOPE TAGGED RECOMBINANT GROWTH ARREST SPECIFIC GENE
TITLE OF INVENTION: PROTEINS
FILE REFERENCE: CEN-5015USNP
CURRENT FILIGATION NUMBER: US/10/671,054
CURRENT FILING DATE: 2003-09-24
PRIOR APPLICATION NUMBER: 60/413157
PRIOR APPLICATION NUMBER: 60/413157
PRIOR APPLICATION NUMBER: 60/413157
PRIOR APPLICATION NUMBER: 5004-09-24
NUMBER OF SEQ ID NOS: 3
SOFTWARE: Patentin version 3.2
SEQ ID NO 3.2
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Publication No. US20040157774A1
GENERAL INFORMATION:
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Best Local Similarity
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TYPE: PRT
ORGANISM: Homo sapiens
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                                                                                      SCHSGFELSSDGRTCQDIDECADSEACGEARCKNLPGSYSCLCDEGFAYSSQEKACRDVD
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                                      FSGTPVIRLRFKRLQPTRLVAEFDFRTFDPEGILLFAGGHQDSTWIVLALRAGRLELQLR
                                                                                                                                                                SCHSGFELSSDGRTCQDIDECADSEACGEARCKNLPGSYSCLCDEGFAYSSQEKACRDVD
                   FSGTPVIRLRFKRLQPTRLVAEFDFRTFDPEGILLFAGGHQDSTWIVLALRAGRLELQLR
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100.0%; Pred. No. 7.9e-289;
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GENERAL INFORMATION:
APPLICANT: Lorens, James B.
APPLICANT: Archison, Robert E.
APPLICANT: Friera, Anabella
APPLICANT: Holland, Sacha
APPLICANT: Holland, Sacha
APPLICANT: Holland, Sacha
ITITE OF INVENTION: Modulators of Angiogenesis and
FILE REFERENCE: 021044-005820US
CURRENT APPLICATION NUMBER: US/10/696,909A
CURRENT FILING DATE: 2003-10-29
PRIOR APPLICATION NUMBER: US 60/512,251
PRIOR APPLICATION NUMBER: US 60/421,989
PRIOR FILING DATE: 2002-10-29
NUMBER OF SEQ ID NOS: 72
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 41
LENGTH: 678
TYPE: PRT
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Matches 678; Conservative
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OTHER INFORMATION:
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SCHSGFELSSDGRTCQDIDECADSEACGEARCKNLPGSYSCLCDEGFAYSSQEKACRDVD
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                                                                        TPNPCDRKGTQACQDLMGNFFCLCKAGWGGRLCDKDVNECSQENGGCLQICHNKPGSFHC
                                                                                         TPNPCDRKGTQACQDLMGNFFCLCKAGWGGRLCDKDVNECSQENGGCLQICHNKPGSFHC
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100.0%;
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Pred. No. 7.9e-289;
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; TYPE: PRT
; ORGANISM: Homo Sapiens
US-10-756-149-4949
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US-10-756-149-4949
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Publication No. US20050181375A1

GENERAL INFORMATICN:
APPLICANT: Aziz, Natasha
APPLICANT: Zlotnik, Albert

TITLE OF INVENTION: MCTHODS OF DIAGNOSIS OF METASTATIC CANCER, COMPOSITIONS OF INVENTION: MCTHODS OF SCREENING FOR MODULATORS OF METASTATIC CANCER

FILE REFERENCE: file
CURRENT APPLICATION NUMBER: US/10/756,149

CURRENT FILING DATE: 2004-01-12

NUMBER OF SEQ ID NOS: 5818

SOFTWARE: Patentin version 3.2

SEQ ID NO 4949

LENGTH: 678
                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 678; Conserv
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                          SCHSGFELSSDGRTCQDIDECADSEACGEARCKNLPGSYSCLCDEGFAYSSQEKACRDVD
                                                                                                    TPNPCDRKGTQACQDLMGNFFCLCKAGWGGRLCDKDVNECSQENGGCLQICHNKPGSFHC
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         SCHSGFELSSDGRTCQDIDECADSEACGEARCKNLPGSYSCLCDEGFAYSSQEKACRDVD
                                                                             TENECDRKGTQACQDLMGNFFCLCKAGWGGRLCDKDVNECSQENGGCLQICHNKEGSFHC
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100.0%; Pr
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APPLICANT: YE, WELLON COMPOSITIONS AND METHODS FOR THE DIAGNOSIS AND TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DIAGNOSIS AND TITLE OF INVENTION: TREATMENT OF DISORDERS INVOLVING ANGIOGENESIS FILE REFERENCE: P3235P1C10

CURRENT APPLICATION NUMBER: US/10/223,085

CURRENT FILING DATE: 2002-08-16

PRIOR APPLICATION NUMBER: US 60/213,637

PRIOR FILING DATE: 2000-06-23

PRIOR APPLICATION NUMBER: US 60/213,637

PRIOR APPLICATION NUMBER: US 60/219,556

PRIOR APPLICATION NUMBER: US 60/219,556

PRIOR APPLICATION NUMBER: US 60/220,624

PRIOR APPLICATION NUMBER: US 60/220,664

PRIOR FILING DATE: 2000-07-25

PRIOR PILING DATE: 2000-07-25

PRIOR APPLICATION NUMBER: US 60/220,664

PRIOR PILING DATE: 2000-07-25

PRIOR APPLICATION NUMBER: US 60/220,664

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PRIOR APPLICATION NUMBER: US 60/220,664

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PRIOR APPLICATION NUMBER: US 60/220,664

PRIOR FILING DATE: 2000-07-25

PRIOR APPLICATION NUMBER: US 60/220,664

PRIOR FILING DATE: 2000-07-25

PRIOR FILING DATE: 2000-07-25

PRIOR APPLICATION NUMBER: US 60/222,695

PRIOR FILING DATE: 2000-08-02
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US-10-223-085-48
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Publication No. US20030100497A1
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APPLICANT: Ferral
APPLICANT: Gerbe
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Gurney, Austin L.
Hillan, Kenneth J.
Marsters, Scot A.
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Gerber, Hanspeter
Geritaen, Mary E.
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Wood, William I
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kevin P.
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PRIOR APPLICATION NUMBER: US 09/643,657
PRIOR FILING DATE: 2000-08-17
PRIOR APPLICATION NUMBER: PCT/US00/23522
PRIOR APPLICATION NUMBER: PCT/US00/23522
PRIOR APPLICATION NUMBER: PCT/US00/23328
PRIOR FILING DATE: 2000-08-24
PREMAINING PRIOR APPLICATION APPLICATION data removed - UMBER OF SEQ ID NOS: 383
SEQ ID NO 48
LENGTH: 673
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Best Local Simi
Matches 547;
          Sequence 48, Application US/10223084 Publication No. US20030105011A1 GENERAL INFORMATION:
APPLICANT: Baker, Kevin
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81.6%; Pred. No. 1.4e-233;
tive 45; Mismatches 75;
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APPLICANT: SCEPHAN, GEAL-FRIATPPE F.
APPLICANT: Williams, P.Mickey
APPLICATION COMPOSITIONS AND METHODS FOR THE DIAGNOSIS AND
TITLE OF INVENTION. TREATMENT OF DISORDERS INVOLVING ANGIOGENESIS
FILE REFERENCE: 93235P1C5
FILE REFERENCE: 9335P1C5
FILE REFERENCE: 9325P1C5
FILE REFERENCE: 9335P1C5
FILE REFERENCE: 9325P1C5
FILE REFERENCE: 9325P
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Best Local Similarity
Matches 547; Conserv
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IRLREXRLQPTRLVAEFDFRTFDPEGILLFAGGHQDSTWIVLALRAGRLELQLRYNGVGR
                                                                                  CEQTCVNSPGSYTCHCDGRGGLKLSPDMDTCEDILFCVPFSMAKSVKSLYLGRMFSGTPV
                                                                                                                 CEQVCVNSPGSYTCHCDGRGGLKLSQDMDTCEDILPCVPFSVAKSVKSLYLGRMFSGTPV
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Gerber, Hanspeter
General Mary E.
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Gurney, Austin L.
Hillan, Kenneth J.
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Pred. No. 1.46
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APPLICANT: Baker, Kevin P.
APPLICANT: Gerritsen, Mary E.
APPLICANT: Gerritsen, Mary E.
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APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Austin L.
APPLICANT: Goddard, Austin L.
APPLICANT: Goddard, Austin L.
APPLICANT: Hillan, Kenneth J.
APPLICANT: Marsters, Scot A.
APPLICANT: Wetphan, Jean-Philippe F.
APPLICANT: Williams p. Mickey
APPLICANT: Wetphan, Jean-Philippe F.
APPLICANT: Wetphan, Jean-Philippe F.
APPLICANT: Williams p. Mickey
APPLICANTON: TREATMENT OF DISORDERS INVOLVING ANGIOGENESIS
FILE REFERENCE: P9235PIC6
CURRENT APPLICATION NUMBER: US/10/223,088
CURRENT APPLICATION NUMBER: US 10/081,056
PRIOR APPLICATION NUMBER: US 60/213,637
PRIOR APPLICATION NUMBER: US 60/213,637
PRIOR APPLICATION NUMBER: US 60/220,624
PRIOR APPLICATION NUMBER: US 60/220,625
PRIOR APPLICATION NUMBER: US 60/220,625
PRIOR APPLICATION NUMBER: US 60/220,624
PRIOR APPLICATION NUMBER: US 60/220,625
PRIOR APPLICATION NUM
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US-10-223-088-48
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                                                   RESULT 8
US-10-223-090-48
US-10-223-090-48
Sequence 48, Application US/10223090
Publication No. US20030105013A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Ferrara, Napoleone
APPLICANT: Gerritsen, Mary E.
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, Audrey
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; SEQ ID NO 48
; LENGTH: 673
; LENGTH: 673
; TYPE: PRT
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Gerritsen, Mary E.
Goddard, Audrey
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Godowski, Paul J.
Gurney, Austin L.
Hillan, Kenneth J
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SEQ ID NO 48
LENGTH: 673
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DIAGNOSIS AND TITLE OF INVENTION: TREATMENT OF DISORDERS INVOLVING ANGIOGENESIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: P3235P1C2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 60/219,556
FILING DATE: 2000-07-20
APPLICATION NUMBER: US 60/220,624
FILING DATE: 2000-07-25
APPLICATION NUMBER: US 60/220,664
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APPLICATION NUMBER: PCT/US00/20710
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                                          PRIOR FILING DATE: 2000-06-23
PRIOR APPLICATION NUMBER: US 60/219,556
PRIOR APPLICATION NUMBER: US 60/220,624
PRIOR APPLICATION NUMBER: US 60/220,664
PRIOR FILING DATE: 2000-07-25
PRIOR FILING DATE: 2000-07-25
PRIOR PRIOR APPLICATION NUMBER: US 60/220,664
PRIOR PRIOR APPLICATION NUMBER: US 60/220,664
PRIOR PILING DATE: 2000-07-28
PRIOR PILING DATE: 2000-08-02
PRIOR PILING DATE: 2000-08-02
PRIOR PILING DATE: 2000-08-02
PRIOR PILING DATE: 2000-08-17
PRIOR PILING DATE: 2000-08-24
PRIOR APPLICATION NUMBER: US 60/230,978
PRIOR PILING DATE: 2000-09-24
PRIOR PILING DATE: 2000-09-24
PRIOR PILING DATE: 2000-09-07
PRIOR PILING DATE: 2000-09-07
PRIOR APPLICATION NUMBER: US 60/230,978
PRIOR PILING DATE: 2000-09-07
PRIOR APPLICATION NUMBER: US 60/232,887
PRIOR APPLICATION NUMBER: US 60/232,887
PRIOR PILING DATE: 2000-09-18
PRIOR PILING DATE: 2000-09-18
PRIOR PILING DATE: 2000-09-18
PRIOR PILING DATE: 2000-09-18
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APPLICANT: Ye, Weilan
APPLICANT: Ye, Weilan
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DIAGNOSIS AND
TITLE OF INVENTION: TREATMENT OF DISORDERS INVOLVING ANGIOGENESIS
FILE REFERENCE: P3235P1C4
CURRENT APPLICATION NUMBER: US/10/223,087
CURRENT APPLICATION NUMBER: US 10/081,056
PRIOR APPLICATION NUMBER: US 00/213,637
PRIOR APPLICATION NUMBER: US 60/213,637
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                                                                                         OR APPLICATION NUMBER: PCT/US00/23522
OR FILING DATE: 2000-08-23
OR APPLICATION NUMBER: PCT/US00/23328
OR APPLICATION NUMBER: PCT/US00/23328
OR APPLICATION NUMBER: US 60/230,978
OR APPLICATION NUMBER: US 60/232,887
APPLICATION NUMBER: US 09/664,610 FILING DATE: 2000-09-18 APPLICATION NUMBER: US 09/665,350 FILING DATE: 2000-09-18
                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: PCT/US00/20710 FILING DATE: 2000-07-28
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Godowski, Paul J.
Gurney, Austin L.
Hillan, Kenneth J.
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Best Local
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APPLICATION NUMBER: PCT/US01/21066
APPLICATION 1001-06-29
TITING DATE: 2001-06-29
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APPLICATION NUMBER: US 09/866,034
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FILING DATE: 2000-12-20
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APPLICATION NUMBER: US 09/816,744
FILING DATE: 2001-03-22
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FILING DATE: 2000-10-24
APPLICATION NUMBER: US 09/709,238
FILING DATE: 2000-11-08
APPLICATION NUMBER: PCT/US00/30952
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APPLICATION NUMBER: PCT/US00/32678
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NUMBER: US 09/747,259
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SHSCPPVEHA
                                                                      RLAVLERHLRSPVLTFAGGLPDVPVTSAPVTAFYRGCMTLEVNRRLLDLDEAAYKHSDIT
                                                                                                                               KLKKQLVVLAVEHTALALMEIKVCDGQEHVVTVSLRDGEATLEVDGTRGQSEVSAAQLQE
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                                                      RLDTLKTHLQGSVHTYVGGLPEVSVISAPVTAFYRGCMTLEVNGKILDLDTASYKHSDIT
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APPLICANT: Ye, Weilian
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DIAGNOSIS AND
TITLE OF INVENTION: TREATMENT OF DISORDERS INVOLVING ANGIOGENESIS
FILE REFERENCE: P3235D1C8
CURRENT FILLING DATE: 2002-08-16
PRIOR APPLICATION NUMBER: US/10/223,083
CURRENT FILLING DATE: 2002-08-16
PRIOR APPLICATION NUMBER: US 60/213,637
PRIOR FILLING DATE: 2002-02-20
PRIOR APPLICATION NUMBER: US 60/213,637
PRIOR FILLING DATE: 2000-06-23
PRIOR APPLICATION NUMBER: US 60/219,556
PRIOR APPLICATION NUMBER: US 60/219,556
PRIOR APPLICATION NUMBER: US 60/219,556
PRIOR FILLING DATE: 2000-07-20
PRIOR FILLING DATE: 2000-07-25
PRIOR APPLICATION NUMBER: US 60/220,664 US-10-223-083-48 GENERAL INFORMATIO
APPLICANT: Baker,
APPLICANT: Ferra Sequence 48, Application US/10223083 Publication No. US20030119112A1 APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT:
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APPLICANT: APPLICANT: INFORMATION Ferrara, Naputer Gerber, Hanspeter Geritsen, Mary E Watanabe, Colin K. Wood, William I. Stephan, Jean-Philippe Watanabe, Colin K. Pan, Godowski, Paul J. Gurney, Austin L. Hillan, Kenneth J. Goddard, Audrey Godowski, Paul Williams, P.Mickey Marsters, Scot James Kevin P Þ יד<u>י</u>

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PRIOR FILING DATE: 2000-07-28
PRIOR PPLICATION NUMBER: US 60/222,695
PRIOR PPLICATION NUMBER: US 60/222,695
PRIOR PPLICATION NUMBER: US 09/643,657
PRIOR APPLICATION NUMBER: US 09/643,657
PRIOR APPLICATION NUMBER: PCT/US00/23522
PRIOR FILING DATE: 2000-08-23
PRIOR FILING DATE: 2000-08-23
PRIOR APPLICATION NUMBER: PCT/US00/23328
PRIOR FILING DATE: 2000-08-24
Remaining Prior Application data removed -
NUMBER OF EQ ID NOS: 383
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LENGTH: 673
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                          AHSCPPVEPA 676
                                                        KLKKQLVVLAVEHTALALMEIKVCDGQEHVVTVSLRDGEATLEVDGTRGQSEVSAAQLQE
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SHSCPPVEHA
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Pred. No. 1.4e
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RESULT 11

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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DIAGNOSIS AND TITLE OF INVENTION: TREATMENT OF DISORDERS INVOLVING ANGIOGENESIS FULL REFERENCE: P3235PIC9
CURRENT APPLICATION NUMBER: US/10/223,089
CURRENT FILING DATE: 2002-08-16
PRIOR APPLICATION NUMBER: US 10/081,056
PRIOR FILING DATE: 2002-02-20
PRIOR APPLICATION NUMBER: US 60/213,637
PRIOR APPLICATION NUMBER: US 60/213,637
PRIOR APPLICATION NUMBER: US 60/213,637
PRIOR PILING DATE: 2000-06-23
PRIOR PILING DATE: 2000-07-26
PRIOR PILING DATE: 2000-07-25
PRIOR APPLICATION NUMBER: US 60/220,624
PRIOR APPLICATION NUMBER: US 60/220,664
PRIOR APPLICATION NUMBER: US 60/220,664
PRIOR PILING DATE: 2000-07-25
PRIOR APPLICATION NUMBER: US 60/220,664
PRIOR PILING DATE: 2000-07-25
PRIOR APPLICATION NUMBER: US 60/220,664
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PRIOR APPLICATION NUMBER: US 60/220,664
PRIOR PILING DATE: 2000-07-25
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PRIOR FILING DATE: 2000-07-28
PRIOR PRILING DATE: 2000-08-02
PRIOR PILING DATE: 2000-08-02
PRIOR PILING DATE: 2000-08-02
PRIOR APPLICATION NUMBER: US 09/643,657
PRIOR APPLICATION NUMBER: PCT/US00/23522
PRIOR APPLICATION NUMBER: PCT/US00/23328
PRIOR APPLICATION NUMBER: PCT/US00/23328
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 48
LENGTH: 673
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Best Local Similarity
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NUMBER OF SEQ ID NOS: 383
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ORGANISM: Murine
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Gerber, Hanspeter
Gertsen, Mary E
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Godowski, Paul J.
Gurney, Austin L.
Hillan, Kenneth J.
                                                             ELSSDGRTCQDIDECADSEACGEARCKNLPGSYSCLCDEGFAYSSQEKACRDVDECLQGR
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CEQVCVNSPGSYTCHCDGRGGLKLSQDMDTCEDILPCVPFSVAKSVKSLYLGRMFSGTPV
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Watanabe, Colin K.
Wood, William I.
Williams, P.Mickey
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Pred. No. 1.4e
45; Mismatches
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APPLICANT: YE. HEALTS.
TITLE OF INVENTION: COMPOSITIONS AND METHODS
TITLE OF INVENTION: TREATMENT OF DISORDERS
FILE REFERENCE: 923591C7
CURRENT APPLICATION NUMBER: US/10/223,081
CURRENT FILING DATE: 2002-08-16
PRIOR APPLICATION NUMBER: US 10/081,056
PRIOR FILING DATE: 2002-06-23
PRIOR APPLICATION NUMBER: US 60/213,637
PRIOR FILING DATE: 2000-06-23
PRIOR FILING DATE: 2000-07-20
PRIOR PPLICATION NUMBER: US 60/219,556
PRIOR FILING DATE: 2000-07-26
PRIOR PPLICATION NUMBER: US 60/220,624
PRIOR APPLICATION NUMBER: US 60/220,664
PRIOR APPLICATION NUMBER: US 60/220,664
PRIOR FILING DATE: 2000-07-25
PRIOR APPLICATION NUMBER: US 60/220,664
PRIOR PILING DATE: 2000-07-26
PRIOR PILING DATE: 2000-07-26
PRIOR APPLICATION NUMBER: US 60/222,695
PRIOR APPLICATION NUMBER: US 60/223,657
PRIOR FILING DATE: 2000-08-02
PRIOR FILING DATE: 2000-08-07
PRIOR FILING DATE: 2000-08-17
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US-10-223-081-48
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APPLICANT: Ferrarz
APPLICANT: Gerrier
APPLICANT: Gerrier
APPLICANT: Goddway
APPLICANT: Godows)
APPLICANT: Hillan
APPLICANT: Hillan
APPLICANT: Marste:
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APPLICANT: Watanal
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Gerber, Hanspeter
Corritsen, Mary E.
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Godowski, Paul J.
Gurney, Austin L.
Hillan, Kenneth J.
Marsters, Scot A.
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Stephan, Jean-Philippe
Watanabe, Colin K.
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Ye, Weilan
                                                                                                                                                                                                                                                                                                                                                                                             Wood, William I.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      671
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RESULT 13
US-10-223-082-48
US-10-223-082-48
; Sequence 48, Application US/10223082
; Publication No. US20030191059A1
; GENERAL INFORMATION:
; APPLICANT: Basker, Kevin P.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Gerber, Hanspeter
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PRIOR FILING DATE: 2000-08-23
PRIOR APPLICATION NUMBER: PCT/US00/23328
PRIOR FILING DATE: 2000-08-24
Remaining Prior Application data removed
NUMBER OF SEQ ID NOS: 383
SEQ ID NO 48
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TYPE: PRT
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NUMBER OF SEQ ID NOS: 383
SEQ ID NO 48
LENGTH: 673
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Best Local Similarity
Matches 547; Conserv
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PRIOR EILING DATE: 2002-02-20
PRIOR APPLICATION NUMBER: US 60/213,637
PRIOR FILING DATE: 2000-06-23
PRIOR APPLICATION NUMBER: US 60/219,556
PRIOR FILING DATE: 2000-07-20
PRIOR APPLICATION NUMBER: US 60/220,624
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CURRENT APPLICATION NUMBER: US/10/223,082
CURRENT FILING DATE: 2002-08-16
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DIAGNOSIS AND
TITLE OF INVENTION: TREATMENT OF DISORDERS INVOLVING ANGIOGENESIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 2000-08-23
PRIOR APPLICATION NUMBER: PCT/US00/23328
PRIOR FILING DATE: 2000-08-24
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R PILING DATE: 2000-07-25

RAPPLICATION NUMBER: US 60/220,664

RAPPLICATION NUMBER: US 60/220,664

RAPPLING DATE: 2000-07-25

APPLING DATE: 2000-07-25
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APPLICATION NUMBER: US 60/222,695
FILING DATE: 2000-08-02
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APPLICATION NUMBER: PCT/US00/20710
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                                                              SLASDGQTCQDIDECTDSDTCGDARCKNLPGSYSCLCDEGYTYSSKEKTCQDVDECQQDR
                                                                                                                                                                                                       KKGTHICQDLMGNFFCVCTDGWGGRLCDKDVNECVQKNGGCSQVCHNKPGSFQCACHSGF
                                                                                                                                                                                                                                                                                               EELCSREEAREVFENDPETDYFYPRYLDCINKYGSPYTKNSGFATCVQNLPDQCTPNPCD
              IRLRFKRLQPTRLVAEFDFRTFDPEGILLFAGGHQDSTWIVLALRAGRLELQLRYNGVGR
                                                                                                                                                                                                                                                                           EEVCSKEEAREVFENDPETEYFYPRYQECMRKYGRPEEKNPDFAKCVQNLPDQCTPNPCD
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IRLRFKRLOPTRILAEFDFRTFDPEGVLFFAGGRSDSTWIVLGLRAGRLELQLRYNGVGR
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Watanabe, Colin K.
Wood, William I.
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Gurney, Austin L.
Hillan, Kenneth J.
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81.6%; Pred. No. 1.4e-233;
tive 45; Mismatches 75;
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APPLICANT: Paoni, N. F.
APPLICANT: Stephan, J-P F.
APPLICANT: Watenabe, C.K.
APPLICANT: Wood, W.I.
APPLICANT: Williams, P.M.
APPLICANT: Ye, Weilan
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DIAGNOSIS AND TITLE OF INVENTION: TREATMENT OF DISORDERS INVOLVING ANGIOGENESIS FILE REFERENCE: P3235R1C1
CURRENT APPLICATION NUMBER: US/10/305,654
CURRENT APPLICATION NUMBER: US/10/305,654
NUMBER OF SEQ ID NOS: 383
SEQ ID NO 48
LENGTH.
                                                                                                                                                                                                                           ; LENGTH: 673
; TYPE: PRT
; ORGANISM: Murine
US-10-305-654-48
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
APPLICANT:
                                                                                                                                                                                  Query Match
Best Local Similarity
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Godowski, Paul J.
Gurney, Austin L.
Hillan, Kenneth J.
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                                                          EEVCSKEEAREVFENDPETEYFYPRYQECMRKYGRPEEKNPDFAKCVQNLPDQCTPNPCD
                                                                       BELCSREEAREVFENDPETDYFYPRYLDCINKYGSPYTKNSGFATCVQNLPDQCTENPCD
                                                                                                                PGPAA-ALGTALLLLLASESSHTVLLRAREAAQFLRPRQRRAYQVFEEAKQGHLERECV
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No. US20030224984A1
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                                                                                                                                                                       Conservative
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                                                                                                                                                                                   80.4%;
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                                                                                                                                                                                     Score 2955.5;
Pred. No. 1.46
                                                                                                                                                                       Mismatches
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RESULT 15
US-10-081-056-48
CURRENT APPLICATION NUMBER: US/10/081,056
CURRENT FILING DATE: 2002-02-02
PRIOR APPLICATION NUMBER: PCT/US01/21735
PRIOR FILING DATE: 2001-07-09
PRIOR APPLICATION NUMBER: US 60/219,556
PRIOR FILING DATE: 2000-07-20
PRIOR APPLICATION NUMBER: US 60/220,624
PRIOR FILING DATE: 2000-07-25
PRIOR APPLICATION NUMBER: US 60/220,664
PRIOR APPLICATION NUMBER: PCT/US00/20710
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Publication No. US20040043927A1
                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                       APPLICANT: Ye, Weilan
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DIAGNOSIS AND
TITLE OF INVENTION: TREATMENT OF DISORDERS INVOLVING ANGIOGENESIS
FILE REFERENCE: P3235P1C1
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Ye, Weilan
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Paoni, Nicholas F.
Stephan, Jean-Philippe
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Gerber, Hanspeter
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Gurney, Austin L.
Hillan, Kenneth J
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Marsters, Scot A.
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; LENGTH: 673
; TYPE: PRT
; ORGANISM: Murine
US-10-081-056-48
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SEQ ID NO 48
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PRIOR APPLICATION NUMBER: PCT/US01/00000
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OR APPLICATION NUMBER: US 09/796,498
OR FILING DATE: 2001-02-28
OR APPLICATION NUMBER: PCT/US01/06520
OR FILING DATE: 2001-03-01
OR APPLICATION NUMBER: US 09/802,706
OR APPLICATION NUMBER: US 09/802,706
OR APPLICATION NUMBER: US 09/808,689
OR APPLICATION NUMBER: US 09/808,689
OR FILING DATE: 2001-03-14
OR APPLICATION NUMBER: US 09/808,689
OR FILING DATE: 2001-03-14
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OR FILING DATE: 2000-12-01

OR APPLICATION NUMBER: US 09/747,259

OR FILING DATE: 2000-12-20

OR APPLICATION NUMBER: PCT/US00/34956

OR FILING DATE: 2000-12-20

OR APPLICATION NUMBER: US 09/767,609
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OR APPLICATION NUMBER: US 60/242,922
OR FILING DATE: 2000-10-24
OR APPLICATION NUMBER: US 09/709,238
OR FILING DATE: 2000-11-08
OR APPLICATION NUMBER: PCT/US00/30952
OR APPLICATION NUMBER: PCT/US00/30873
OR FILING DATE: 2000-11-08
OR FILING DATE: 2000-11-10
OR FILING DATE: 2000-11-10
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FILING DATE: 2001-03-22
APPLICATION NUMBER: US 09/828,366
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APPLICATION NUMBER: PCT/US01/17443
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FILING DATE: 2001-05-25
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APPLICATION NUMBER: US 09/664,610
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APPLICATION NUMBER: US 09/643,657
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2: /EMC_Celerra_SIDS3/ptodata/2/iaa/6_COMB.pep:*

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ALIGNMENTS

; TOPOLOGY: unknown ; MOLECULE TYPE: protein US-08-282-141-2 US-08-282-141-2 Sequence 2, Application US/08282141 Patent No. 5538861 FILING DATE: CLASSIFICATION: 435 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: GENERAL INFORMATION: CORRESPONDENCE ADDRESS: ADDRESSE: Amgen Inc. STREET: 1840 Dehavillanc CITY: Thousand Oaks STATE: California COUNTRY: United States ZIP: 91320-1789 APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: COMPUTER READABLE FORM: MEDIUM TYPE: Plopy disk COMPUTER: IBM PC compactible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/282,141 APPLICANT: Brancolini, Claudio APPLICANT: Manfioletti, Guidalberto TITLE OF INVENTION: Stimulating Fact NUMBER OF SEQUENCES: LENGTH: 678 amino acids TYPE: amino acid STRANDEDNESS: unknown h 98.7%; Score 3628; DB 1; 1 Similarity 100.0%; Pred. No. 1.5e-309; 78; Conservative 0; Mismatches 0; E: Amgen Inc. 1840 Dehavilland Schneider, Claudio Varnum, Brian Avanzi, Giancarlo Claudio 2 Drive Factor for the AXL Receptor Length 678;

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APPLICANT: Mather, Jennie P.

APPLICANT: Li, Ronghao

APPLICANT: Chen, Jian

TITLE OF INVENTION: ISOLATING AU

NUMBER OF SEQUENCES: 9

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                      Sequence 2, Application Patent No. 5714385
          APPLICATION NUMBER: US/08/
FILING DATE: 10-MAY-1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Lee, Wendy M.
                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: 5.25 inch, 360 Kb i
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                       ADDRESSEE: Genentech, Inc.
STREET: 460 Point San Bruno B
CITY: South San Francisco
STATE: California
COUNTRY: USA
ZIP: 94080
  REGISTRATION NUMBER:
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RESULT 3
US-08-435-436-2
US-08-08-2721139
; Sequence 2, Application US/08435436
; Patent No. 5721139
; GENERAL INFORMATION:
; APPLICANT: Mather, Jennie P.
; APPLICANT: Li, Ronghao
; APPLICANT: Chen, Jian
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Best Local S
Matches 678
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-1994
TELEFAX: 415/952-9881
TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 678 amino acid
TYPE: amino acid
TOPOLOGY: linear
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                  AAQLQERLAVLERHLRSPVLTFAGGLPDVPVTSAPVTAFYRGCMTLEVNRRLLDLDEAAY
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Best Local Similarity
Matches 678; Conserv
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SEQUENCE CHARACTERISTICS:
LENGTH: 678 amino acids
TYPE: amino acid
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REGISTRATION NUMBER: 00,000
REFERENCE/DOCKET NUMBER: 940
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/00 FILING DATE: 10-MAY-1995 CLASSIFICATION: 424 PRIOR APPLICATION DATA:
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CURRENT APPLICATION DATA:
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MEDIUM TYPE: 5.25 inch,
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ADDRESSEE: Genentech
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NUMBER OF SEQUENCES:
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TELEFAX: 415/952-9881
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STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
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OPERATING SYSTEM: PC-DOS/MS-DOS
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PGSGFAFYSLDYMRTPLDVGTESTWEVEVVAHIRPAADTGVLFALWAPDLRAVPLSVALV
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Pred. No. 1.5e-309;
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US-08-438-863-2
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                                                                                                                                                                                                                                                                                                              NAME: Lee, Wendy M.

REGISTRATION NUMBER: 00,000

REFERENCE/DOCKET NUMBER: 946

TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-1994

TELEPAX: 415/952-9881

TELEPAX: 415/952-9881

TELEPAX: 415/952-9881

INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
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Matches
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 94080
COMPUTER READABLE FORM:
MEDIUM TYPE: 5.25 inch, 360 Kb floppy
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS

COMPUTER: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/0
FILING DATE: 10-MAY-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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SOFTWARE: patin (Genen
CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech,
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                                                                                                                                                                                                                                                            LENGTH: 678 amino TYPE: amino acid TOPOLOGY: linear
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CITY: South San Francisco
STATE: California
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121 TPNPCDRKGTQACQDLMGNFFCLCKAGWGGRLCDKDVNECSQENGGCLQICHNKPGSFHC 180
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                                                                                                     MAPSLSPGPAALRRAPQLLLLLLAAECALAALLPAREATQFLRPRQRRAFQVFEEAKQGH
                                 LERECVEELCSREEAREVFENDPETDYFYPRYLDCINKYGSPYTKNSGPATCVQNLPDQC
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100.0%; Pr
100.0%; 0;
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                                                                                                                                                                         Score 3628; DB 1;
Pred. No. 1.5e-309;
0; Mismatches 0;
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RESULT 5
US-08-438-864-2
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                                               APPLICANT: Chen, Jian
APPLICANT: Godowski, Paul J.
APPLICANT: Mark, Melanie R.
APPLICANT: Mark, Melanie P.
APPLICANT: Mark, Melanie P.
APPLICANT: Mather, Jennie P.
APPLICANT: Li, Ronghao
TITLE OF INVENTION: RSE RECEPTOR ACTIVATION
NUMBER OF SEQUENCES: 2
CORRESSORDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STATE: OSA
CITY: South San Francisco
STATE: OSA
COUNTRY: USA
ZIP: 94080
COMPUTER READABLE FORM:
MEDIUM TYPE: 5.25 inch, 360 Kb floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: 5.25 inch, 360 Kb floppy disk
COMPUTER: 1BM PC compatible
OPERATION SYSTEM: PC-DOS/MS-DOS
SOFTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/438,864
FILING DATE: 10-MAY-1995
CLASSIFICATION: 435
PRIOR ADDITION TATA:
APPLICATION ATA:
                                                                                                                                                                                                                                                                                                                                                                                     Sequence 2, Applicati
Patent No. 5955420
GENERAL INFORMATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/402253
FILING DATE: 10-MAR-1995
ATTORNEY/AGENT INFORMATION:
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REGISTRATION NUMBER: 00,000
REFERENCE/DOCKET NUMBER: 929P1.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-1994
TELEFAX: 415/952-9881
TELEFAX: 415/952-9881
TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 6678 amino acids
TYPE: amino acids
TYPE: amino acids
TYPE: amino acids
RESULT 6
US-08-438-862-2
; Sequence 2, Application U
; Patent No. 6033660
; GENERAL INFORMATION:
; APPLICANT: Mather, Je
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Query Match 98.7%; Score 3628; DB 1; 1
Best Local Similarity 100.0%; Pred. No. 1.5e-309;
Matches 678; Conservative 0; Mismatches 0;
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                                                                                                                             DYHSTKKLKKQLVVLAVEHTALALMEIKVCDGQEHVVTVSLRDGEATLEVDGTRGQSEVS
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AAQLQERLAVLERHLRSPVLTFAGGLPDVPVTSAPVTAFYRGCMTLEVNRRLLDLDEAAY
                         AAQLQERLAVLERHLRSPVLTFAGGLPDVPVTSAPVTAFYRGCMTLEVNRRLLDLDEAAY
                                                                                              DYHSTKKLKKQLVVLAVEHTALALMEIKVCDGQEHVVTVSLRDGEATLEVDGTRGQSEVS
                                                                                                                                                                                        PGSGFAFYSLDYMRTPLDVGTESTWEVEVVAHIRPAADTGVLFALWAPDLRAVPLSVALV
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660

540 480 480

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360 300 300 240 240 180 180 120 120 60

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US-08-438-862-2
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 678 amin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Lee, Wendy M.
REGISTRATION NUMBER: 00,000
REFERENCE/DOCKET NUMBER: 946-
TELECOMPUNICATION INFORMATION:
TELEPHONE: 415/25-1994
TELEFAX: 415/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Li, Ronghao
APPLICANT: Chen, Jian
TITLE OF INVENTION: ISOI
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: 5.25 inch, 360 Kb f
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid TOPOLOGY: linear
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STREET: 460 Point San Bruno
CITY: South San Francisco
STATE: California
   421
                                   421
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                   LTVGGIPFHEKDLVQPINPRLDGCMRSWNWLNGEDTTIQETVKVNTRMQCFSVTERGSFY
                                                                                                                                                                                                                           ECLQGRCEQVCVNSPGSYTCHCDGRGGLKLSQDMDTCEDILFCVFFSVAKSVKSLYLGRM 300
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                                                                                      YNGVGRVTSSGPVINHGMWQTISVEELARNLVIKVNRDAVMKIAVAGDLFQPERGLYHLN
                                                                                                                                                                                                     ECLQGRCEQVCVNSPGSYTCHCDGRGGLKLSQDMDTCEDILPCVPFSVAKSVKSLYLGRM
                                                                                                                                    FSGTPVIRLRFKRLQPTRLVAEFDFRTFDPEGILLFAGGHQDSTWIVLALRAGRLELQLR
                                                                                                                                                            FSGTPVIRLRFKRLQPTRLVAEFDFRTFDPEGILLFAGGHQDSTWIVLALRAGRLELQLR
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LTVGGIPFHEKDLVQPINPRLDGCMRSWNWLNGEDTTIQETVKVNTRMQCFSVTERGSFY
                                                                  YNGVGRVTSSGPVINHGMWQTISVEELARNLVIKVNRDAVMKIAVAGDLFQPERGLYHLN
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Conservative 0;
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Pred. No. 1.5e-309;
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; TOPOLOGY: US-08-402-253-2
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Patent No. 6211142
GENERAL INFORMATION:
APPLICANT: Chen, Jian
APPLICANT: Hammonds, R. Glenn
                                                                                                                                           Matches
                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                         TELEFAX: 415/952-9881
TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 678 amino acids
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NAME: Wendy M. Lee
REGISTRATION NUMBER: 00,000
REFERENCE/DOCKET NUMBER: 929
TELECOMMUNICATION INFORMATION:
TELECHIONE: 415/225-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/402,253
FILING DATE: 10-MAR-1995
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESS:
ADDRESSE: GenenceN, Inc.
STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTMARE: patin (Genentech) CURRENT APPLICATION DATA:
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MEDIUM TYPE: 5.25 inch,
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APPLICANT: Godowski, Paul J.
APPLICANT: Mark, Melanie R.
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                                                                                                                                                                                                                                                                                                                                 TELEPHONE: 415/22-9881
                                                                                                                                                                                                                                                          TYPE:
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                                                                                                                                   98.7%; Score 3628; DB 2; 1
100.0%; Pred. No. 1.5e-309;
cive 0; Mismatches 0;
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Sequence 2, Application US/08443866B
; Patent No. 6255068
; GENERAL INFORMATION:
APPLICANT: Godowski, Paul J.
APPLICANT: Hammonds, R. Glenn
APPLICANT: Hammonds, R. Glenn
APPLICANT: HAMEN, Melanie R.
TITLE OF INVENTION: RSE RECEPTOR ACTIVATION
NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STATE: California
COUNTRY: USA
ZIP: 94080
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Wilpatin (Genentech)
CURRENT APPLICATION NUMBER: US/08/443,866B
FILING DATE: 31-May-1995
CLASSIFICATION NUMBER: 08/402253
FRIOR APPLICATION NUMBER: 08/402253
FRIOR APPLICATION NUMBER: 08/402253
FRIING DATE: 10-MAR-1995
ATTORNEY/AGENT INFORMATION:
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US-08-443-866B-2
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               RESULT 9
US-08-628-747-2
US-08-628-747-2
; Sequence 2, Application U
; Patent No. 6169070
; Patent INFORMATION:
; GENERAL INFORMATION:
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REGISTRATION NUMBER: 00,000
REFERENCE/DOCKET NUMBER: P092
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-1994
TELEPAX: 415/952-9881
TELEFAX: 415/952-9881
TELEFX: 910/371-7168
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LEMENT: CHARACTERISTICS:
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Best Local Similarity
Matches 678; Conserv
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TYPOLOGY: Line-
          APPLICANT:
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100.0%; Pred. No. 1.5e-309;
                                                    US/08628747
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; MOLECULE TYPE: protein
US-08-628-747-2
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Best Local S
Matches 677
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APPLICATION NUMBER: US/08/628,747
FILING DATE: 17-APR-1996
CLASSIFICATION DATA:
APPLICATION UNWBER: US 08/402,253
PRIOR APPLICATION DATA:
APPLICATIO
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 678 amino acid
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APPLICANT: Hammonds, R. Glenn
APPLICANT: Mark, Melanie
APPLICANT: Mark, Melanie P.
APPLICANT: Li, Ronghao
TITLE OF INVENTION: RECEPTOR ACTIVATION BY
NUMBER OF SEQUENCES: 30
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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3T: One DNA Way
: South San Francisco
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                                  FSGTPVIRLRFKRLQPTRLVAEFDFRTFDPEGILLFAGGHQDSTWIVLALRAGRLELQLR
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US-08-282-141-3
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Patent No. 5538861
GENERAL INFORMATION:
APPLICANT: Schneider, Claudio
APPLICANT: Varnum, Brian
APPLICANT: Avanzi, Giancarlo
                                                                                                                            Query Match
Best Local Similarity
                                                                                                               Matches
                                                                                                                                                                                                                                                                                    CLASSIFICATION: 435
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                   ZIP: 91320-1789
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compartible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS
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APPLICANT: Manfioletti, Guidalberto
TITLE OF INVENTION: Stimulating Factor for the
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CITY: Thousand Oaks
                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE:
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                                                                                                          Score 2955.5; DB 1;
Pred. No. 1.7e-250;
5; Mismatches 75;
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RESULT 11
US-08-435-434-1
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APPLICANT: Chen, Jian
TITLE OF INVENTION: ISOLATING AFTITLE OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ANDRESSEE: Genentech, Inc.
                                                                                                                                                                                                                                                                                                                                                 Sequence 1, Applic Patent No. 5714385
                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Mather
APPLICATION NUMBER: US/08/
FILING DATE: 10-MAY-1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: 5.25 inch, 360 Kb i
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                         STREET: 460 Point San Bruno
CITY: South San Francisco
STATE: California
COUNTRY: USA
ZIP: 94080
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REGISTRATION NUMBER: 946-
REFERENCE/DOCKET NUMBER: 946-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-1994
TELEPAX: 415/952-9881
TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 673 amino acids
TYPE: amino acids
TYPE: amino acid
TOPOLOGY: linear
                US-08-435-436-1
; Sequence 1, Application
; Patent No. 5721139
; GENERAL INFORMATION:
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Best Local Similarity
Matches 547; Conserv
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                                                                                                             SHSCPPVEHA
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81.6%;
      Jennie
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Pred. No. 1.7e-250;
5; Mismatches 75;
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US-08-435-436-1
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TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 673 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/435,436
FILLING DATE: 10-MAY-1995
CLASSIFICATION: 424
PRIOR APPLICATION NUMBER:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY JACENT INFORMATION:
NAME: Lee, Wendy M.
REGISTRATION NUMBER: 00,000
REFERENCE/DOCKET NUMBER: 946-:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
STREET: 460 Point San Bruno
CITY: South San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Li, Ronghao
APPLICANT: Chen, Jian
TITLE OF INVENTION: ISOLATING AND CULTURING SCHWANN CELLS
NUMBER OF SEQUENCES: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 5.25 inch,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
TOPOLOGY: linear
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PFKESELVQPINPRLDGCMRSWNWLNGEDSAIQETVKANTKMQCFSVTERGSFFPGNGFA
                                                               ITSSGPTINHGMWQTISVEELERNLVIKVNKDAVMKIAVAGELFQLERGLYHLNLTVGGI
                                                                                                                                 IRLRFKRLQFTRLLAEFDFRTFDFEGVLFFAGGRSDSTWIVLGLRAGRLELQLRYNGVGR
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81.6%; Pred. No. 1.7e-250;
45: Mismatches 75;
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US-08-438-863-1
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                                                                                                                                                      Query Match
Best Local Similarity
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Patent No. 5849585
GENERAL INFORMATION:
APPLICANT: Mather, Jennie P.
APPLICANT: Ronghao Li
                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                             TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Lee, Wendy M.
REGISTRATION UNMBER: 00,000
REFERENCE/DOCKET NUMBER: 946
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-1994
TELEFAX: 415/952-9881
                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: PC-DOS/MS
SOFTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: 5.25 inch, 360 Kb f
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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TITLE OF INVENTION:
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                                                                                                                                                                                                                                            TYPE: amino acid
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                                                                                                                                                                                                                           TOPOLOGY: linear
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                EELCSREEAREVFENDPETDYFYPRYLDCINKYGSPYTKNSGFATCVQNLPDQCTFNPCD 126
                                                                                  PGPAALRRAPQLLLLLLAAECALAALLPAREATQFLRPRQRRAFQVFEEAKQGHLERECV
EEVCSKEEAREVFENDPETEYFYPRYQECMRKYGRPEEKNPDFAKCVQNLPDQCTPNPCD
                                                                   PGPAA-ALGTALLLLLASESSHTVLLRAREAAQFLRPRQRRAYQVFEEAKQGHLERECV
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460 Point San Bruno
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Pred. No. 1.7e-250;
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RESULT 14
US-08-438-864-1
; Sequence 1, Application
; Patent No. 5955420
; Patent INFORMATION:
                                                                                                                                                                                               APPLICANT: Chen, Jian
APPLICANT: Hammond, Paul J.
APPLICANT: Godowski, Paul J.
APPLICANT: Mark, Melanie R.
APPLICANT: Mark, Melanie P.
APPLICANT: Mather, Jennie P.
APPLICANT: Li, Ronghao
TITLE OF INVENTION: RSE RECEPTOR ACTIVATION
NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESS:
ADDRESSE: Genentech, Inc.
STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STATE: California
COUNTRY: USA
                                            COMPUTER READABLE FORM:
MEDIUM TYPE: 5.25 inch, 360 Kb fl
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/438,864
FILING DATE: 10-MAY-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                 CLASSIFICATION:
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US-08-438-862-1 ; Sequence 1, Application US/08438862 ; Patent No. 6033660

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AUTORNEY/AGENT INFORMATION:
NAME: Wendy M. Lee
REGISTRATION NUMBER: 00,000
REFERENCE/DOCKET NUMBER: 929P:
TELECOMMUNICATION INFORMATION:
TELEPAX: 415/252-1994
TELEFAX: 415/952-9881
TELEFAX: 910/371-7168
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 673 amino acids
TYPE: amino acid
TYPE: amino acid
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                         AHSCPPVEPA 676
                                                                 RLAVLERHLRSPVLTFAGGLPDVPVTSAPVTAFYRGCMTLEVNRRLLDLDEAAYKHSDIT
                                                                                                     KLKKQLVVLAVEHTALALMEIKVCDGQEHVVTVSLRDGEATLEVDGTRGQSEVSAAQLQE
                                                                                                                                                                                                                                                              ITSSGPTINHGMWQTISVEELERNLVIKVNKDAVMKIAVAGELFQLERGLYHLNLTVGGI
SHSCPPVEHA
                                                  FYSLOYMRTPLDVGTESTWEVEVVAHIRPAADTGVLFALWAPDLRAVPLSVALVDYHSTK
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Pred. No. 1.7e-250;
5; Mismatches 75;
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TELEPAX: 415/952-9881
; TELEX: 910/371-7168
; INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 673 amino acids
TYPE: amino acid
TYPE: amino acid
TYPE: linear
US-08-438-862-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 80.4%; Score 2955.5; DB 2; Length 673; Best Local Similarity 81.6%; Pred. No. 1.7e-250; Matches 547; Conservative 45; Mismatches 75; Indels 3;
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MEDIUM TYPE: 5.25 inch, 360 K
COMPUTER: IBM PC comparible
COMPUTER: IBM PC comparible
OPERATING SYSTEM: PC-DOS/MS-D
SOFTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
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FILING DATE: 10-MAX-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Mather, Jennie P. APPLICANT: Li, Ronghao APPLICANT: Chen, Jian
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REGISTRATION NUMBER: 00,000
REFERENCE/DOCKET NUMBER: 94
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CITY: South San Francisco
STATE: California
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                                 CEQVCVNSPGSYTCHCDGRGGLKLSQDMDTCEDILPCVPFSVAKSVKSLYLGRMFSGTPV 306
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PFHEKDLVQPINPRLDGCMRSWNWLNGEDTTIQETVKVNTRMQCFSVTERGSFYPGSGFA
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                                                                 RLAVLERHLRSPVLTFAGGLPDVPVTSAPVTAFYRGCMTLEVNRRLLDLDEAAYKHSDIT
                                                                                                 KLKKQLVVLAVEDVALALMEIKVCDSQEHTVTVSLREGEATLEVDGTKGQSEVSTAQLQE
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Search completed: June 29, 2006, 11:17:34 Job time : 53 secs

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Sequence 42, Appl Sequence 21, Appl Sequence 21, Appl Sequence 371, Appl Sequence 375, App Sequence 2765, Appl Sequence 2, Appl Sequence 2, Appl Sequence 2, Appl Sequence 5, Appl Sequence 5, Appl Sequence 3, Appl Sequence 31, Appl Sequence 38, Appl Sequence 39, Appl Sequence 4910, App Sequence 4810, App Sequence 3263, App Sequence 3263, App Sequence 3263, Appl Sequence 3263, Appl Sequence 3263, App Sequence 3263, App Sequence 38, Appl Sequence 3263, App Sequence 3263, App Sequence 3263, App Sequence 3263, App Sequence 18, Appl

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length: 2000000000
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| MMC_Celerra_SIDS3/ptodata/1/pubpaa/US09_NEW_PUB.pep:*
| EMC_Celerra_SIDS3/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
| MMC_Celerra_SIDS3/ptodata/1/pubpaa/US07_NEW_PUB.pep:*
| MMC_Celerra_SIDS3/ptodata/1/pubpaa/US08_NEW_PUB.pep:*
| MMC_Celerra_SIDS3/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
| MMC_Celerra_SIDS3/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
| MMC_Celerra_SIDS3/ptodata/1/pubpaa/US11_NEW_PUB.pep:*
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Listing first 45 summaries
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    US-10-505-928-451
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US-11-267-942-3
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US-11-217-997-38
US-10-511-937-2962
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US-11-183-218-10
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Sequence 451, Appli Sequence 57, Appli Sequence 77, Appli Sequence 4, Appli Sequence 4, Appli Sequence 2, Appli Sequence 2, Appli Sequence 3062, Appli Sequence 10, Appl Sequence 10, Appli Sequence 10, Appli Sequence 11, Appli Sequence 28, Appli Sequence 28, Appli Sequence 14, Appli Sequence 14, Appli Sequence 14, Appli Sequence 27, Appli Sequence 28, Appli Sequence 27, Ap
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CURRENT APPLICATION NUMBER: US/10/505,928
CURRENT FILING DATE: 2004-08-27
PRIOR APPLICATION NUMBER: US 60/363,019
PRIOR FILING DATE: 2002-03-07
NUMBER OF SEQ ID NOS: 866
SOFTWARE: Patentin 3.2
SEQ ID NO 451
LENGTH: 1821
CRGANISM: Homo sapiens
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US-10-505-928-451
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                                                                     SODMDTCEDILPC 283
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US-11-175-714-4
US-11-196-749-29
US-11-175-714-4
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US-11-175-714-8
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US-11-267-942-5
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                                                                             ; TYPE: PRT; ORGANISM: Homo sapiens US-10-505-928-676
                                                                                                                                                                                                                                                                                                                                            Sequence 676, Application US/10505928 Publication No. US20060088532A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn version 3.3 SEQ ID NO 5
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                                                                                                                                                      SOFTWARE: PatentIn 3.2 SEQ ID NO 676
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Best Local Similarity
                   Query Match
Best Local Similarity
     Matches
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APPLICANT: Albig, Allan R.
TITLE OF INVENTION: Fibulin-3 and Uses Thereof
FILE REFERENCE: 2879-109
                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/505,928
CURRENT FILING DATE: 2004-08-27
PRIOR APPLICATION NUMBER: US 60/363,019
PRIOR FILING DATE: 2002-03-07
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PRIOR FILING DATE: 2004-11-04
PRIOR APPLICATION NUMBER: 60/687,129
PRIOR FILING DATE: 2005-06-03
                                                                                                                                                                                                                                                                                     APPLICANT: Ludwig Institute for Cancer Research et al. TITLE OF INVENTION: LYMPHATIC ENDOTHELIAL GENES FILE REFERENCE: 28967/39178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 19
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CURRENT FILING DATE: 2005-11-04
                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 866
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ORGANISM: Mus musculus
                                                                                                                                      LENGTH: 493
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   65;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EACRG--DMMCVNQNGGYLCIPRTNPVYRGPYSNPYSTSYSGPYPAAAPPVPASNYPTIS 108
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No. US20060094054A1
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   Conservative
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8.7%; Score 321; DB 6
38.2%; Pred. No. 3e-18;
Live 30; Mismatches
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Pred. No. 1.6e-18;
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                                     DB 6; Length 493;
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APPLICANT: Schiemann, William P.
APPLICANT: Albig, Allan R.
TITLE OF INVENTION: Fibulin-3 and Uses Thereof
FILE REFERENCE: 2879-109
CURRENT APPLICATION NUMBER: US/11/267,942
CURRENT FILING DATE: 2005-11-04
PRIOR APPLICATION NUMBER: 60/625,598
PRIOR FILING DATE: 2004-11-04
PRIOR APPLICATION NUMBER: 60/625,129
PRIOR APPLICATION NUMBER: 60/687,129
PRIOR APPLICATION NUMBER: 60/687,129
PRIOR FILING DATE: 2005-06-03
NUMBER OF SEQ ID NOS: 19
SOFTWARE: PatentIn version 3.3
SEQ ID NO 3
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US-11-267-942-2
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; LENGTH: 493
; TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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Publication No. US20060094054A1
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CURRENT APPLICATION NUMBER: US/11/267,942

CURRENT FILING DATE: 2005-11-04

PRIOR APPLICATION NUMBER: 60/625,598

PRIOR FILING DATE: 2004-11-04

PRIOR APPLICATION NUMBER: 60/687,129

PRIOR APPLICATION NUMBER: 60/687,129

PRIOR PILING DATE: 2005-06-03
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APPLICANT: Albig, Allan R.
TITLE OF INVENTION: Fibulin-3 and Uses Thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 19
SOFTWARE: PatentIn version 3.3
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LENGTH: 493
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            233 GSFYCQCSPGFQLAANNYTCVDINECDASNQCAQ-QCYNILGSFICQCNQGYELSSDRLN
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US-10-196-749-408
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US-10-196-749-408
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   Best Loc
Matches
                                                                                                                                                 SEQ ID NO 408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
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CURRENT FILING DATE: 2002-07-16
PRIOR APPLICATION NUMBER: 10/052586
PRIOR FILING DATE: 2002-01-15
PRIOR APPLICATION NUMBER: 60/059263
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REPERENCE: P3430R1C340
                                                                                                                                                                Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 612
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Baker, Kevin P.
                                                                                           ORGANISM: Homo
                                                                                                                              LENGTH: 448
                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                          APPLICATION NUMBER: 60/063541
                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 60/063540 FILING DATE: 1997-10-28
                                                                                                                                                                                                                                                                                                                  FILING DATE: 1997-10-21
                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 60/063486
                                                                                                                                                                                                                                                                                                                                                                                                                                          FILLING DATE: 1997-09-18
APPLICATION NUMBER: 60/059266
FILING DATE: 1997-09-18
APPLICATION NUMBER: 60/062250
                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 60/063121
                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 1997-10-17
APPLICATION NUMBER: 60/063120
                                                                                                                                                                                                                        APPLICATION NUMBER: 60/063544
                                                                                                                                                                                                                                         FILING DATE: 1997-10-28
                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 1997-10-24
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     66;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GSFYCQCNPGFQLAANNYTCVDINECDASNQCAQ-QCYNILGSFICQCNQGYELSSDRLN 291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CRDVDECLQGR--CEQVCVNSPGSYTCHCDGRGGLKLSQDMDTCEDILPC 283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Smith, Victoria Watanabe, Colin K. Wood, William I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Zhang, Zemin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gurney, Austin L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Goddard, Audrey
     Conservative
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                                                                                           Sapien
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                                                                                                                                                                                                        1997-10-28
                   8.7%;
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38.8%; Pred. No. 3.6e-18;
rative 30; Mismatches 64
     23;
 Score 319; DB
Pred. No. 3.8e
23; Mismatches
                 DB 6;
3.8e-18;
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                                    Length 448;
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 14;
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Gaps
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US-11-267-942-4
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                                                                                                                                                                                                                                                                                                                                              Sequence 38, Application US/11217997 Publication No. US20060111561A1 GENERAL INFORMATION:
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                                                                                                                  APPLICANT:
                                                                                                                                                                                                                               APPLICANT: Valerie L. Gerlach
APPLICANT: Elma R. Fernandes
APPLICANT: Richard A. Shimket
APPLICANT: Meera Patturajan
APPLICANT: Vladimir Y. Gusev
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Schiemann, William P.
APPLICANT: Albig, Allan R.
TITLE OF INVENTION: Fibulin-3 and Uses Thereof
FILE REFERENCE: 2879-109
CURRENT APPLICATION NUMBER: US/11/267,942
CURRENT FILING DATE: 2005-11-04
CURRENT APPLICATION NUMBER: US/11/217,997
                          TITLE OF INVENTION: NOVEL PROTEINS AND NUCLEIC ACIDS ENCODED THEREBY FILE REFERENCE: Cura 551 CIP
                                                                      APPLICANT:
                                                                                                                                                                                      APPLICANT:
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SOFTWARE: PatentIn version 3.3
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PRIOR FILING DATE: 2004-11-04
PRIOR APPLICATION NUMBER: 60/687,129
PRIOR FILING DATE: 2005-06-03
                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 493
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       184 VPGSYSCTCNPGFTLNEDGRSCQDVNECATENPCVQT-CVNTYGSLICRCDPGYELEEDG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CEDIDECRISSYLCOYOCVNEPGKFSCMCP--OGYEVVRSR-TCQDINEC 338
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GSFYCQCSPGFQLAANNYTCVDINECDASNQCAQ-QCYNILGSFICQCNQGYELSSDRLN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GSFHCSCHSGFELSSDGRTCQDIDECADSEACGEARCKNLPGSYSCLCDEGFAYSSQEKA 235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DECTSGTHNCRTDQVCINLRGSFTCQCLPGYQKRGEQC-VDIDECTVP-PYCHQRCVNTP 232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             VHCSDMDECSFSEFLCQHECVNQPGTYFCSCP--PGYILLDDNRSCQDINEC 292
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DECATDSHQCNPTQICINTEGGYTCSCTDGYWLLEGQCL---DIDEC--RYGYCQQLCAN
                                                                 Muralidhara Padigaru
                                                                                                                                                                                   Stacie (Casman) Navara
Velizar T. Tchernev
                                                                                                                                       David W. Anderson
Xiaojia (Sasha) Guo
                                                                                                                                                                                                                                                                             Elma R. Fernandes
Richard A. Shimkets
                                                                                                                  Luca Rastelli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               65; Indels
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Sequence 2, Application US/10766760

Publication No. US20060127915A1

GENERAL INFORMATION:

APPLICANT: Klein, Michael

APPLICANT: Schultz, Dennis

TITLE OF INVENTION: Gene Mutation Associated with

TITLE OF INVENTION: Age-Related Macular Degeneration

FILE REFERENCE: 49321-117

CURRENT APPLICATION NUMBER: US/10/766,760

CURRENT FILING DATE: 2004-01-27

PRIOR APPLICATION NUMBER: 60/443,214

PRIOR FILING DATE: 2003-01-27

NUMBER OF SEQ ID NOS: 186

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 2

LENGTH: 5635
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US-10-766-760-2
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SEQ ID NO 38
LENGTH: 1418
TYPE: PRT
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ORGANISM: Homo sapiens
-11-217-997-38
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 2003-06-03
APPLICATION NUMBER: 10/055,877
FILING DATE: 2002-01-22
APPLICATION NUMBER: 60/262,892
FILING DATE: 2001-01-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 60/264,478 FILING DATE: 2001-01-26 APPLICATION NUMBER: 60/263,351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 2001-01-24
APPLICATION NUMBER: 60/264,117
FILING DATE: 2001-01-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 60/263,598 FILING DATE: 2001-01-23 APPLICATION NUMBER: 60/263,799
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 2001-03-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 60/272,870
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 60/264,139 FILING DATE: 2001-01-25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        264 QVVRGLARCECHVGYQLAADGKACEDVDECAAGLAQC----AHGCLNTQGSFKCVCHAGY 319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    59 GHLERECVEELCSR------EEAREVFENDPETDYFYPRYLDCINKYGSPYTKN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ----GGRLCDK----DVNECSQENGGCLQICHNKPGSFHCSCHSGFELSSDGRTCQDIDE 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 S---GFATCVQNL------PDQCTPN--PCDRKGTQACQDLMGNFFCLCKAGW 148
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GGCQHHCVQLTITRHRCQCRPGFQLQEDGRHCVRRSP-----CANRNGSCMHRC 263
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8.6%; Score 316; DB 7; Length 1418;
29.8%; Pred. No. 3.1e-17;
ative 29; Mismatches 97; Indels 5
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APPLICANT: ROSENberg, Steven
TITLE OF INVENTION: METHODS AND COMPOSITIONS FC
TITLE OF INVENTION: AND MONITORING TRANSPLANT
FILE REFERENCE: 506612000104
CURRENT APPLICATION NUMBER: US/10/511,937
CURRENT FILING DATE: 2004-10-19
PRIOR APPLICATION NUMBER: PCT/US2003/012946
PRIOR FILING DATE: 2003-04-24
PRIOR FILING DATE: 2003-04-24
PRIOR APPLICATION NUMBER: US 10/131,831
PRIOR APPLICATION NUMBER: US 10/325,899
PRIOR APPLICATION NUMBER: US 10/325,899
PRIOR FILING DATE: 2002-12-20
NUMBER OF SEQ ID NOS: 3117
SOFTWARE: PatentIn version 3.2
SEQ ID NO 2962
LENGTH: 444
TYPER: PRT
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US-10-511-937-2962
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                        Matches
                                                                                                                                                                           Query Match
Best Local
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Local Similarity 29.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5128 YY-----CSCPKGLTIAADGRTCQDIDECALGRHTCHAGQDCDNTIGSYRCVVRCGSGF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               206 ACGEARCKNIPGSYSC--LCDEGFAYSSQEKACRDVDECLQG----RCEQVCVNSPGSYT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          102 PYTKNSGFATCVQNLP------DQCTPNPCDRKGTQACQDLMGNFFCLCKAGWGG 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      50 FQVFEEAKQGHLEREC-----VEELCSREEAREVFENDPETDYFYPRYLDCINKYGS
  64
                                74 EAREVFENDPETDYFYPRYLDCINKYGSPYTKNSGFATCVQNLPDQCTPNPCDRKGTQAC
                                                                                                             14 RAPQLILILILAAECALAALLPAREATQFLRPRQRRAFQVFEEAKQGHLERECVEELCSRE
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                                                                                                                                                        66;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          T: Rosenberg, Steven
INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING
                                                                            4 QALRILCILLGLQGCLAAVFVTQEEAHGVLHRRRRANAFLEELRPGSLERECKEEQCSFE
                                                                                                                                                                             Similarity
  EAREIFKDAERTKLFWISYSD-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CHCD----GRGGLKLSQDMDTCEDI-LPC
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Prentice, James
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                                                                                                                                                        8.1%; Score 296; DB 6;
35.5%; Pred. No. 2.7e-16;
ative 27; Mismatches 65
                                                                                                                                                      27;
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Pred. No. 3.8e-15;
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                                                                                                                                                                                               Length 444;
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US-11-183-218-8
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                                   Sequence 8, Application US/11183218
Publication No. US20060088906A1
GENERAL INFORMATION:
APPLICANT: Neose Technologies, Inc.
APPLICANT: DeFrees, Shawn
APPLICANT: Zopf, David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 8.1%; Score 296; DB 6; I Best Local Similarity 35.5%; Pred. No. 2.7e-16; Matches 66; Conservative 27; Mismatches 65;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn version 3.2 SEQ ID NO 3014
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 3014, Application US/10511937 Publication No. US20060088836A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 10/325,899
PRIOR FILING DATE: 2002-04-24
PRIOR APPLICATION NUMBER: US 10/325,899
PRIOR FILING DATE: 2002-12-20
                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 2002-12-20 NUMBER OF SEQ ID NOS: 3117
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CURRENT FILING DATE: 2004-10-19
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TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING
TITLE OF INVENTION: AND MONITORING TRANSPLANT REJECTION
FILE REFERENCE: 506612000104
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Fry, Kirk
Woodward, Robert
Bayer, Robert
Hakes, David
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Publication No.
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APPLICANT:
APPLICANT:
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Best Local
TITLE OF INVENTION: ERYTHROPOIETIN: REMODELING AND TITLE OF INVENTION: GLYCOCONJUGATION OF ERYTHROPOIETIN FILE REFERENCE: 040853-01-5083-US02
CURRENT APPLICATION NUMBER: US/11/183,218
CURRENT FILING DATE: 2005-07-15
                                                                                                                                                            APPLICANT: Neose Technologies,
APPLICANT: DeFrees, Shawn
APPLICANT: Zopf, David
APPLICANT: Bayer, Robert
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PRIOR FILING DATE: 2003-04-09
PRIOR APPLICATION NUMBER: PCT/US02/32263
PRIOR FILING DATE: 2002-10-09
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CURRENT FILING DATE: 2005-07-15
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ORGANISM: Homo sapiens
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OR APPLICATION NUMBER: US 60/387,292
OR FILING DATE: 2002-06-07
OR APPLICATION NUMBER: US 60/344,692
OR FILING DATE: 2001-11-19
OR APPLICATION NUMBER: US 60/334,301
OR FILING DATE: 2001-11-28
OR APPLICATION NUMBER: US 60/334,233
OR FILING DATE: 2001-11-28
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FILLING DATE: 2002-08-16
APPLICATION NUMBER: US 60/396,594
FILLING DATE: 2002-07-17
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Bayer, Robert
Hakes, David
Chen, Xi
Bowe, Caryne
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Pred. No. 2.7e-16;
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Best Local Similarity
Matches 65; Conserv
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SEQ ID NO 10
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                                                                                                                                                                                                                                                                                                                     Sequence 18, Appropriate Publication No.
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CURRENT APPLICATION NUMBER: US/11/217,997
CURRENT FILING DATE: 2005-08-31
PRIOR APPLICATION NUMBER: 10/453,372
PRIOR FILING DATE: 2003-06-03
                                                             APPLICANT: Muralidhara Padigaru
TITLE OF INVENTION: NOVEL PROTEINS AND NUCLEIC ACIDS
FILE REFERENCE: Cura 551 CIP
                                                                                                                                                                                                                                                                                    APPLICANT: Valerie L.
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APPLICATION NUMBER: US 60/334,233
FILING DATE: 2001-11-28
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FILING DATE: 2002-08-15
APPLICATION NUMBER: US 60/396,594
FILING DATE: 2002-07-17
FILING DATE: 2002-07-17
FILING DATE: 2002-07-17
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                                                                                                                                               Meera Patturajan
Vladimir Y. Gusev
Stacie (Casman) Navara
Velizar T. Tchernev
David W. Anderson
Xiaojia (Sasha) Guo
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No. US20060111561A1
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Richard A. Shimkets
                                                                                                                   Mei Zhong
                                                                                                                                     Luca Rastelli
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Pred. No. 8.7e-15;

    See File Wrapper or PALM

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Sequence 8, Application US/11264243

Publication No. US20060134121A1

GENERAL INFORMATION:
APPLICANT: Thurston, Gavin
APPLICANT: Moguera, Irene
TITLE OF INVENTION: DL14 Antagonists, As
TITLE OF INVENTION: Methods Thereof
FILE REFERENCE: 2070A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/272,870 PRIOR FILING DATE: 2001-03-02
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PRIOR FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
CURRENT APPLICATION NUMBER: US/11/264,243
CURRENT FILING DATE: 2005-10-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT ORGANISM: Homo sapiens
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FILING DATE: 2001-01-25
APPLICATION NUMBER: 60/264,478
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 60/263,799 FILING DATE: 2001-01-24
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                                                                                                                                                                                                                                                                                                                                                                          228 AYSSQEKACRDVDECLQG--RCEQVCVNSPGSYTCHC-----DGRGGLKLSQD-MDT
                                                                                                                                                                                                                                                                                                                                                                                                             215 HHCVQLTITRHRCQCRPGFQLQEDGRHCVRRSPCANRNGSCMHRCQVVRGLARCECHVGY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   65 --CVEEL-----CSREEAREVFENDPETDYFYPRYLDCINKYGSPYTKN 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4 LEEARAAGRAVVLALVLLLLPAVPVGASVPPRP----LLPLQPGMPHVCAEQELTLVGRR
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; PRIOR APPLICATION NUMBER: 60/623,658
; PRIOR FILLING DATE: 2004-10-29
; NUMBER OF SEQ ID NOS: 17
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 8
; LENGTH: 2003
; TYPE: PRT
; ORGANISM: Homo sapien
US-11-264-243-8
Search completed: June 29, 2006, 11:31:56 Job time : 22 secs
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Best Local Similarity 31.3%; Pred. No. 9e-14;
Matches 76; Conservative 21; Mismatches 77; Indels 69; Gaps 12;
                                                                                                                               427 PTCHQDLDECLMAQQGPSPCEHGGSCLNTPGSFNCLCPPGYTGSRCEADHNECLS-QPCH 485
                                                                                                                                                             233 EKACRDVDECLQGR-----CEQ--VCVNSPGSYTCHC-DGRGGLKLSQDMDTCEDILPCV 284
                                                                                                                                                                                              188 -----LSSDGRT---CQDIDECADSEACGEARCKNLP--GSYSCLCDEGFAYSSQ 232
370 CIDRVGSFSCLCPPGRTGLLCHLEDMCLSQPCHGDAQCSTNPLTGSTLCLCQPGY---SG 426
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Result
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3 FLAG pept
2 Flag octa
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| ABP60044 ABP960041 ABU11922 ABJ38456 ABU38454 ADA03349 ABG74451 ABU08631 ABU08631 ABU081552 ABU62120 ABB99043 ABU08104 ABU3489 ABU3489 ABU3489 ABU3489 ABU3489 ABU35887 ABU08795 ABU08795 ABU08795 ABU08796 ABU55861 ABP56844 AAE31706 ABP72065 ABU087966 ABU55867 ABU672367 ABD672367 ABD72286 ABU672367 ABD672367 ABD72286 ABB766633 ABU68596 ABU58664 ABU58661 ABP58151 ABP6672367 ABU68596 ABU68596 ABU38697 ABU68596 ABU38693 ABU68644 ABU68696 ABU38696 ABU38696 ABU38696 ABU38696 ABU38696 ABU38696 ABU38696 ABU38696 ABU68634 ABU6863 |
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| MARCH Marc | 47 100.0 8 7 ADM31 47 100.0 8 7 ADL22 47 100.0 8 7 ADL64 47 100.0 8 7 ADM35 47 100.0 8 7 ADM35 47 100.0 8 7 ADM62 47 100.0 8 8 ADG45 47 100.0 8 8 ADG45 47 100.0 8 8 ADG45 |
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RESULT 2
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            by recombinant DNA techniques. The invention also relates to a novel process for producing protein molecules expression vector capable of expressing a heterologous polypeptide in a transformed host cell, the polypeptide consisting of a constituent polypeptide and an attached identification peptide. The identification peptide has an antigenic terminal portion of amino acids. Recombinant DNA techniques are used for the economic production of a desired protein. It is then efficiently purified in a single affinity chromatography step with high yields. The sequence represents an identification peptide of the invention, used in the preparation of a recombinant plasmid containing the human Interleukin 2 gene
The sequence encodes a fusion contruct which links a mutant sequence of human granulocyte-colony stimulating factor to a yeast alpha-factor leader sequence, Fusion proteins capped with this peptide are resistant to intracellular degradation. See also AAN70029, AAN70031-35 and
                                                                                                                                                                                      New pure human granulocyte
by recombinant DNA methods
                                                                                                                                                                                                                                                                                                                                                                                                                           22-APR-1986;
14-NOV-1986;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Granulocyte-colony yeast alpha-factor
                                                                                                                                                                                                                                                             WPI; 1987-300791/43.
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03-OCT-2002
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, Tushinski
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86US-00931458
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leader sequence.
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                                                                                                                                                                                      colony stimulating factor and muteins in high yields, used for potentiating
                                                                                                                    English
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2.1e+06;
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Matches 8
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                                                             The inventors claim a pharmaceutical compsn. which contains an effective amt. of a recombinant human interleukin-3 protein analogue, rhuIL-3, (Asp15,Asp70). The rhuIL-3 analogue has AA SQ in AAR09326. The compsn. may also comprise the N-terminal octapeptide in AAR09327, and a diluent and 1 or more than 1 biological response modifier. The compsn. has a biological specific activity of equal to or more than 4.0 x 10 to the 7 mcg/mg in a human bone marrow proliferation assay, and a binding affinity for human monocyte IL-3 receptors of equal to or more than 4.0 x 10 to the 10 (M to the minus 1). (Updated on 31-OCT-2002 to add missing OS field.) (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                      Compsns. contg. recomincreased biological
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(IL-3) ar
                                            Sequence 8
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31-OCT-2002
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field.)
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                                                                                                                                                                                      5; Page
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ilarity 100.0%;
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                                                                                                                                                                                                                      recombinant non-glycosylated human interleukin-3 gical activity and binding affinity, for treating
                                                                                                                                                                                                                                                                                                   CORP
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Pred. No. 2.1
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thes 0;
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                     Length
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RESULT 4
AAR44496
ID AAR4
RESULT 5
AAR77283
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Matches 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bifunctional recombinant protein - contains particle and analyte binding moieties, used in agglutination assays pref. on whole blood.
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26-MAY-1994
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                                                        Fas ligand; Fas-
self-tolerance;
                                                                                            FLAG peptide
                                                                                                                    25-MAR-2003
05-DEC-1995
                                                                                                                                                        AAR77283;
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             WO9518819-A1
                                   Synthetic.
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                           correct
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                                                        L; cell surface protein; autoimmune disease;
FLAG; purification.
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Pred. No.
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RESULT 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Goodwin
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01-FEB-1994;
Epstein-Barr virus BZLF2 fusion proteins - used for tre immune disease, transplant rejection, allergy, asthma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
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                                                                                                                                                                                       Alderson M, A
Hutt-Fletcher
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                                                                                                               WPI; 1995-393086/50
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UNIV MISSOURI.
US NAT INST OF HEALTH.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      lupus erythematosus; organ transplant rejection; asthma; I cansplant rejection; therapy; cancer; viral disease; mouse;
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                                                                                                                                                                                       Armitage RJ, Cohen
r LM, Spriggs MK;
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94US-00190559.
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Pred. No. 2.1
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RESULT 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    E Barr virus (BBV) protein. To create BZLF2/Fc, this sequence, a mouse interleukin-7 (IL-7) leader sequence (see AAR87021), an immunoglobulin Fc region (see AAR87023) and a flexible linker (see AAR87024) are joined to the extracellular domain (residues 34 to 23) of the BZLF2 sequence (see C AAR87020). BZLF2 proteins are members of the C-type lectin family. The C-C type lectin is capable of binding the beta chain of a major C protein is capable of binding the beta chain of a major C nistocompatibility complex (MRC) class II antigen. Fusion proteins with C antibody formation, proliferation of blood mononuclear cells and C region, can also be created. BZLF2 proteins inhibit antigen. Specific C antibody formation, proliferation of blood mononuclear cells and C cytotoxic T cell responses. They also exhibit superantigen—like activity. The proteins can be used for treating or preventing autoimmune diseases such as myasthenia gravis, multiple sclerosis and systemic lupus carythematosus. Also, for treating organ or tissue transplant rejection and for treating or preventing allergy or asthma. They can be used for treating cancer and viral disease, especially EBV infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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Best Local
                                                                     Polypeptide and modified versions having a receptor tyrosine kinase activity - are expressed in a non:differentiated blood cell but undereduction in the expression level as the non:differentiated cell
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                                                                                                                                                                                                               (ASAH ) ASAHI KASEI
                                                                                                                                                                                                                                                                                                                                                                                                      IBI FLAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                    IBI FLAG epitope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 8 AA;
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                                                                                                                                    1995-215266/28
DB; AAQ92644.
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                                                                                                                                                                                                                                                                                                                                                                                                                  tyrosine kinase; receptor; non-differentiated; blood cell;
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8; Conserv
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                                                                                                                                                                                   Oono
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Pred. No. 2.1e+06;
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                                                                     but undergo
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AAQ92644 encodes AAR76469 the IBI FLAG epitope, used in the prepn.

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Example 8; Page 39;

73pp; Japanese.

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                                                        Query Match
Best Local
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                                                                                                                        This FLAG-epitope tag peptide has been fused to a Fas receptor, which contains a cytoplasmic region binding a novel FADD protein (Fasassociating protein with novel death domain, AAT39397), which modulates apoptosis induced by activation of the receptor by ligand binding. DNA encoding the tag is fused to the Fas gene and mutants by PCR, downstream of a putative Fas signal peptide sequence, in a plasmid pcDNA3 vector, using a 5', FLAG PCR primer (AAT39398). The resulting fusion protein is expressed in Escherichia coli and used in an assay to demonstrate in vivo association of FADD protein and Fas receptor via death domains. FADD DNA may be used in gene therapy, and FADD protein or a corresponding antibody may be used to screen for agents modulating FADD pathway cellular functions and Fas-associated apoptosis, for use in therapy of e.g. AIDS, inflammation, leukaemia, myocardial infarction, degenerative disease, etc
                                                                                                                                                                                                                                                                                                                                                            FADD protein that binds to cytoplasmic region of Fas receptor - for identifying inhibitors of Fas-associated apoptosis useful for treating e.g. AIDS, leukaemia, stroke, etc.
                                                                                                   Sequence 8
                                                                                                                                                                                                                                                                                                                                    Example 1; Page 51; 96pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fas-associating protein with novel death domain; apoptosis; gene antibody; immunoassay; drug screening; diagnostic; AIDS; antiinflammatory; antitumour; cerebroprotective; neuroprotective
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18-MAY-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FLAG-epitope
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Similarity
8; Conserv
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95US-00443982.
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100.0%; Pred. No. 2.
ive 0; Mismatches
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Pred. No. 2.1e+06;
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                                                     2.1e+06;
                                                                      DB 2;
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RESULT 10
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Best Local S
Matches 8
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                                                                                                                                                                                                                                                                                                                                                 An affinity tag (AAR91066) may be linked to interleukin-1 type-3 receptors (see e.g. AAR91064 and AAR91054) produced by recombinant DNA technology. The peptide facilitates purification of the expressed recombinant protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1996-171614/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-SEP-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Interleukin-1 type-3 receptor; IL-1-3R; immune-associated
vector; antibody; therapy; affinity tag.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Affinity tag peptide
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                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 7; 64pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                               Interleukin-1 type 3 receptor proteins immune-associated diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lovenberg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-SEP-1995;
                                                   Interleukin-1 receptor; human; soluble; N-terminal peptide; antigen; epitope; shuIr-IR; monoclonal antibody; bovine mucosal enterokinase; interleukin-1; II-1; immune response; mammal; diagnosis; therapy; regulation; immune disease; inflammatory disease.
                                                                                                                                25-MAR-2003
01-AUG-1996
                                                                                                                                                                 AAR91045;
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           US5492888-A
                                Synthetic.
                                                                                                          Antigenic N-terminal peptide
                                                                                                                                                                                   AAR91045 standard; peptide; 8 AA
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                                                                                                                                                                                                                                                                                     Conservative
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(first entry)
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                                                                                                                                                                                                                                                                                               100.0%;
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                                                                                                           for fusion to shull-1R.
                                                                                                                                                                                                                                                                                             Score 47;
Pred. No.
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                                                                                                                                                                                                                                                                                                           DB 2;
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                                                                                                                                                                                                                                                                                                         Length 8;
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Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-FEB-1988;
13-OCT-1988;
21-DEC-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sequence can be fused to soluble human interleukin-1 receptor (shuIL-IR). By using this sequence, recombinant shuIL-IR can be assayed and purified easily, by using a monoclonal antibody that reversibly binds this epitope. By using bovine mucosal enterokinase, this sequence can be cleaved from the recombinant shuIL-IR. Proteins capped with this peptide may also be resistant to intracellular digestion in E. coli. The shuIL-IR can be used in a method for suppressing interleukin-1 (IL-I) mediated immune responses in a mammal. The method comprises administering an effective amount of shuIL-IR to the mammal, preferably in a dose of 500ng therapy for regulation of immune or inflammatory activities, in contrast to membrane-bound full-length mature IL-1 receptors. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Use of sol. IL-1 receptors to suppress IL-1-mediated immune responses e.g. for treatment of inflammation in mammals, esp. humans.
                                                                                                                                                                               affinity chromatography; immunoassay; monoclonal antibody; fusion protein; cleavage; cattle mucosal enterokinase; LERK-5; cytokine; fusion protein-tyrosine-kinase; receptor; human fetal brain; cloning; elk; hek; protein-tyrosine-kinase; receptor; human fetal brain; cloning; drug delivery; cell targeting; cytostatic; leukaemia; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This sequence represents an antigenic N-terminal peptide epitope. This sequence can be fused to soluble human interleukin-1 receptor (shuIL-II
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Col 8; 22pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-FEB-1996
                                                                                                                                                                                                                                FLAG; affinity tail; immunoaffinity chromatography; purification;
affinity chromatography; immunoassay; monoclonal antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 8
                                                                                                                                                                                                                                                                           FLAG affinity tail peptide.
                                                                                                                                                                                                                                                                                                        13-APR-1996
                                                                                                                                                                                                                                                                                                                                                                    AAR89288 standard;
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                            08-JUL-1994;
                                                          06-JUL-1995;
                                                                                                                                                  Synthetic
(IMMV ) IMMUNEX CORP
                                                                                                                     WO9601839-A1
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89US-00455488
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                              94US-00271948
                                                            95WO-US008534
                                                                                                                                                                                                                                                                                                                                                                    peptide;
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Pred. No. 2.1
D; Mismatches
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RESULT 12
AAR94931
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Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The sequence represents a FLAG affinity tail, which may be attached to e.g. a full-length or soluble LERK-5 cytokine (ARR89287) to facilitate purification. LERK-5 binds elk and hek receptor protein-tyrosine-kinases (members of the eph/elk family), and may be isolated e.g. by expression from a human fetal brain cell cDNA. The FLAG peptide is highly antigenic, and provides an epitope reversibly bound by a specific monoclonal antibody, enabling rapid immunoassay and purification by immunoaffinity chromatography. The sequence is specifically cleaved by cattle mucosal enterokinase at the residue immediately following the Asp-Lys pairing. Purified LERK-5 may be used in binding assays to detect elk- or hekexpressing cells, to measure elk or hek activity, as a drug delivery for elk or hek receptor-specific cell targeting, e.g. for delivery of
                                                             Dimeric complex having affinity for interleukin-4 - comprises IL-4 receptor homodimer or IL-4-IL2 gamma-c receptor heterodimer, used treatment of e.g. eczema, asthma and HIV.
                                                                                                                                                                                                                                                                                                                                       Interleukin-4 receptor; IL-4; interleukin-2 gamma-c receptor; IL-2;
allergic rhinitis; asthma; atopic dermatitis; eczema; HIV; FLAG.
                                                                                                                                                                                                                                                                                                                                                                                  FLAG
                                      Example
                                                                                                                      WPI; 1996-209810/21.
                                                                                                                                               Vanderslice
                                                                                                                                                                                                   07-OCT-1994;
                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA encoding LERK-5 cytokine capable of binding elk and hek - used as carrier to deliver attached (anti-leukaemic) agents to cells bearing or hek, i.e. human leukaemia cell lines JM and LK63.
                                                                                                                                                                                                                              05-OCT-1995;
                                                                                                                                                                                                                                                         18-APR-1996
                                                                                                                                                                                                                                                                                                                                                                                                             23-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR94931 standard; peptide; 8 AA.
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                                   4; Page 21;
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                         BOULDER INC.
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                                                                                                                                              Cox GM;
                                  53pp; English.
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Pred. No.
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FLAG of a

is an octapeptide (AAR94931) that is used to protein using anti-FLAG monoclonal antibody.

A DNA construct was

S

| DYKDDDDK

Query Match Best Local S Matches 8

8 Similarity

Conservative

100

. 0%;

Score 47; DE Pred. No. 2.1 D; Mismatches

BB .1e+06; 2

Length 8;

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AAW18236
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Best Local
                                                                          The FLAG epitope (AAW18236) is utilised in novel constructs for expression of lysosomal enzymes, e.g. human glucocerebrosidase and alpha-L-iduronidase in transgenic plants. The epitope coding sequence is fused in-frame to the C-terminus of the lysosomal enzyme coding sequence (see also AAT71753) in order to facilitate the detection and purification of the gene product (see also AAW18237). It is designed to be a hydrophilic marker peptide situated on a protein surface to facilitate antibody
Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                  Production of enzymatically active (modified) lysosomal enzyme in transgenic plants - useful in treatment of lysosomal storage disorders.
                                                                                                                                                                                                                                                                                                                             Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Radin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       produced that coded for IL-2Rgamma-c FLAG comprising the 254-amino acid extracellular domain of human interleukin-2 receptor gamma chain plus an additional C-terminal cysteine and the FLAG peptide. The fusion was expressed in Sf9 cells using a baculovirus expression system and purified by affinity chromatography. IL-2Rgamma-c FLAG can be linked to interleukin-4 via a polymeric (PEG) spacer. The resulting dimeric complex has affinity for IL-4 and can be used to treat IL-4 mediated disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1997-202248/18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-SEP-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hurler syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lysosomal enzyme; lysosome; transgenic plant; glucocerebrosidase; alpha-L-iduronidase; IUDA; enzyme replacement therapy; Gaucher disease;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (VIRG ) VIRGINIA TECH INTELLECTUAL (CROP-) CROPTECH DEV CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14-SEP-1995;
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                                                                                                                                                                                                                                                                                                                          Page 37; 111pp;
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Pred. No. 2.1e+06;
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RESULT 14
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Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This peptide comprises the FLAG epitope that is recognised by anti-FLAG monoclonal antibodies. A form of novel growth factor LHL (see AAW32479) was generated by PCR that contains the FLAG epitope at its N-terminus and a strep-tag (see AAW32485) at its C-terminus. The construct is designated LHL.seq (see AAW32481), where L is the immunoglobulin binding entity from Peptostreptococcus magnus and H is residues 42-62 of hen egg lysozyme. The FLAG-epitope was added to facilitate the secretion of LHL.seq into the periplasmic space of host cells. However, this was unsuccessful and LHL.seq needed to be purified from total bacterial lysate. As a result of this, the omph signal peptide was not removed, leading to the formation of LHL.seq multimers. LHL.seq has identical activity to that of LHL, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 3; Page 40; 109pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New catalytic antibody precursors - comprising a B-cell surface molecule binding portion which can induce B-cell mitogenesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1997-489572/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Koentgen F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-MAR-1996;
27-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          02-OCT-1997.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9735887-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Catalytic antibody; growth factor; B-cell mitogenesis; LHL; L protein; hen egg lysozyme; FLAG epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FLAG epitope
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW32484;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAW32484 standard; peptide; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (AMRA-) AMRAD OPERATIONS PTY LTD
DNA binding protein; RNA binding protein; amphipathic peptide;
                                                          FLAG epitope
                                                                                                                        12-NOV-1997
                                                                                                                                                                                                                                          AAW00971 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ۳
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      used in novel methods for the generation of catalytic antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
8; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                            рукоррок в
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DYKUDDDDK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                                        (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      96AU-00008951.
97AU-00005375.
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                                                                                                                                                                                                                                       peptide;
                                                                                                                        entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 47;
Pred. No. :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 8;
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  Query Match
Best Local S
Matches 8
23-OCT-1995;
22-DEC-1995;
09-SEP-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         This peptide sequence comprises a FLAG epitope that can be attached to the N-terminus of a nucleic acid binding protein (NABP) such as a leucine zipper bZIP protein or bHLH protein. Claimed NABPs having an appended acidic extension peptide (see AAW00958-65) can regulate the function of a target nucleic acid or gene to which they are bound, and act as potent dominant-negative regulators of gene transcription, cell growth and cell proliferation. They can be used in cancer therapeutics, to treat diseases caused by eukaryotic microorganisms or by viruses, and as tools for drug development, rational drug design, and drug and gene therapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           acidic extension peptide; gene control; gene regulation; transcription; dominant negative protein; cancer; drug therapy; drug design; leucine zipper; FLAG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New nucleic acid binding proteins - having an acidic extension at the amino-terminus, to increase ability transcription, useful e.g. in cancer therapeutics.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 16; Page 58; 144pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1997-145687/13.
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29-MAY-1996;
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                                                                                                                                                  NR4; haemoprotein asthma; therapy; F
                                                                                                                                                                                         FLAG epitope tag
                                                                                                                                                                                                                                           AAW09824;
                                                                                                                                                                                                                                                                    AAW09824 standard;
                                                   23-OCT-1996;
                                                                                                   WO9715663-A1
                                                                                                                            Synthetic.
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8; Conserv
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                                                                                                                                                                                                                   (first entry)
 95AU-00006135.
95AU-00007276.
96AU-00002208.
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96US-0018496P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            96WO-US012590
                                                   96WO-AU000668
                                                                                                                                                      n receptor;
FLAG.
                                                                                                                                                                                                                                                                    peptide; 8 AA
                                                                                                                                                                  interleukin-13 receptor; cytokine; allergy;
                                                                                                                                                                                                                                                                                                                                                                                   Score 47; DB
Pred. No. 2.1
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SERVICES
                                                                                                                                                                                                                                                                                                                                                                                                2.1e+06;
                                                                                                                                                                                                                                                                                                                                                                                                            DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                            Length 8;
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to regulate gene
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RESULT 17
AAW46330
ID AAW46
AC AAW46
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Best Local
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                                                                                                                                                                                                                                                        Methods for engineering
penton base protein. The presents a binding domain of a chimeric ac penton base protein. The present sequence contains an epitope for antibody that is preferably not present in wild type adenoviryus. The penton base protein binds to cell surface receptors called in The integrins not only provide a binding site for the adenoviral phase protein, but also mediate cellular adhesion to the extracelly matrix molecules. The specification describes
                                                                                                                                                                                                                                                                                                                                                                                                      Bruder
                                                                                                                                                                                                               Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Binding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Using PCR, generated
                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GENV-) GENVEC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA encoding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ing PCR, a derivative of novel haemoprotein receptor NR4 cDNA was merated which encoded the interleukin-3 signal peptide (AAW09823) N-terminal FLAG epitope tag (AAW09824) preceding the mature codir gion (Thr27-Pro424) of murine NR4 (see also AAW09921). The PCR proscioned into the mammalian expression variations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1997-259018/23
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                                                                                                                                                                                                                                                                                                                                                                                                         Mcvey DL,
                                                                                                                                                                                                                                                                                   introducing adenovirus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nicola NA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      animal haemopoietin receptor which interacts with 13 - useful to treat asthma, allergy or condition (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                          16; 56pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     binding
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binding domain; cell surface binding sit
ecule; antibody; adenoviral transfer vec
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                                                                                                                                                                                                                                                          therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
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Pred. No.
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                                          denoviral penton extracellular
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RESULT 18
ADM80476
ID ADM800476
AC ADM800
XX ADM800
XX EST 8
KW EST 8
KW GASTr
KW ulcer
XX Ulcer
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Best Local
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                                            The present sequence represents a peptide sequence encoded by an antisense primer of the invention. The sequence incorporates a recognition site to aid in analysis and purification of the CS141 protein. A set of contiguous and partially overlapping EST sequences (AAV63504-15) are designated CS141. The sequences are isolated from a CDNA library made from gastrointestinal tract tumour and normal tissues. The CS141 gene is useful as a marker for gastrointestinal tract disorders. The methods and products can be used in detecting, diagnosing staging, monitoring, prognosticating, preventing, or determining the predisposition to diseases and conditions of the gastrointestinal tract, such as gastrointestinal tract cancers. Barrel's esophagus, gastric
                                                                                                                                                                                                                                                                                             New gastrointestinal tract specific polynucleotides, CS141 - used develop products for the diagnosis and treatment of e.g. cancers, ulcer, gastritis, Crohn's disease, ulcerative colitis or pancreati
                               ulcer, gastritis,
                                                                                                                                                                                                                                                                                                                                                                                                                                           Billing-Medel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EST sequence; CS141; gastrointestinal tract; gastric ulcer; gastritis; leiomyoma; polyps; ulcerative colitis; pantreatitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        adenovirus into a cell in vitro having a particular cell surface binding site. The adenovirus is contacted with a bispecific molecule (e.g. bispecific antibody) comprising a component that selectively binds a binding domain of the penton base protein of the adenovirus and a second component that selectively binds the cell surface binding site. A complex of the adenovirus and the bispecific molecule is formed, and the cell is contacted with it to allow entry of the adenovirus into the cell. The
                                                                                                                                                                                                                                                                 Example
                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-568280/48
                                                                                                                                                                                                                                                                                                                                                                                                      Roberts-Rapp L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       08-OCT-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EST sequence; CS141; gastric ulcer; gastric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-MAR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                          Hayden
                                                                                                                                                                                                                                                                                                                                                                                                                                             PA,
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                                                                                                                                                                                                                                                               67;
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                                 leiomyoma,
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                                                                                                                                                                                                                                                              English.
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                                 polyps, Crohn's
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Pred. No.
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Klass
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thes 0;
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Crohn's
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                                 ulcerative
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or pancreatitis.
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                                                                                                   diagnosing
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AAW64456
ID AAW6
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AAW54166
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                                  RESULT 20
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                                                                                                                  Matches
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Best Local (
                                                                                                                                                                                               This FLAG epitope was introduced into pCI-neo by ligating complementary oligonuclectides into NheI/EcoRI digested vector. An EcoRI fragment from murine clone B1-1, which codes for novel protein targetting to glycogen (PTG) protein (see AAW54164), was cloned at the EcoRI site of the resulting plasmid. The FLAG epitope-tagged PTG construct was transiently transfected into CHO cells over-expressing the insulin receptor. Results showed that PTG can direct PTI localisation to glycogen both in vivo and in vitro. PTG can be used to develop products for increasing glycogen levels in cells and to develop animal models for the study of e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                  Brady MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-AUG-1996;
12-AUG-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Protein targetting to glycogen; PTG; PP1C; mouse; diabetes; animal model; knockout mouse; FLAG epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FLAG-epitope
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                                                                                                                                                                Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                    Disclosure; Page 24; 71pp; English.
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           AAW64456 standard; protein; 8 AA.
                                                                                                                                                                                                                                                                                                                                          Isolated protein targetting to for increasing glycogen levels the study of e.g. diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                        (WARN ) WARNER LAMBERT CO.
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mes 8; Conserv
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8; Conserv
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                                                                      DYKODDDK 8
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                                                                                                                                                                                                                                                                                                                                                                                                                  Printen JA,
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                                                                                                                  Conservative
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97US-0055243P.
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                                                                                                                             100.0%;
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                                                                                                                  <u>,</u>
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Pred. No.
                                                                                                                             Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                        glycogen
in cells
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                                                                                                                   Mismatches
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                                                                                                                            2.1e+06;
                                                                                                                                                                                                                                                                                                                                                      genes - used to develop products and to develop animal models for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 2;
                                                                                                                                           DB 2;
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                                                                                                                                       Length 8;
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RESULT 21
AAW56763
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Best Local (
                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                               included in compositions and diagnostic kits for diagnosis of breast cancer. The presence of CHK in breast tissue can also be determined using other standard methods (e.g. Northern blotting) or by detecting nucleic acid sequences encoding all/a portion of the protein (e.g. using hybridisation probes). Over-expression of the receptor tyrosine kinase ErbB-2 has previously been associated with the development of breast cancer, and CHK specifically interacts with activated ErbB-2, and may function as a negative regulator of ErbB-2 mediated mitogenic signalling. The compositions may also be used to design drugs (e.g. which incorporate CHK analogues with greater biological activity than CHK) and to identify CHK antagonists and agonists for therapeutic use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          epitope which is used in a method of detecting cancer in breast tissue. The method allows diagnosis of breast cancer in mammals, especially humans. It is based on the discovery that a cytoplasmic protein tyrosine kinase, CHK, is expressed in human breast tissue, but not in adjacent tissue. This protein can be used to raise antibodies which can be tissue. This protein can be used to raise antibodies which can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Detecting breast cancer by detecting Csk homologous kinase expression especially in humans and use of Csk homologous kinase in treatment or prophylaxis of breast cancer and for producing medicaments.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Breast; cancer; matK; CSK homologous kinase; CHK; detection; diagnosis; cytoplasmic protein; tyrosine kinase; ErbB-2; negative regulator; mitogenic signalling.
                                                                                                                                                                                                                                                                                                                                                                    Sequence 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This sequence represents a CSK homologous kinase (CHK) derived flag epitope which is used in a method of detecting cancer in breast tis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1998-399149/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Avraham H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              08-JAN-1997;
16-JUN-1997;
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Tumour necrosis factor related apoptosis ligand;
                                    FLAG peptide epitope used for producing soluble TRAIL polypeptide.
                                                                      05-AUG-1998
                                                                                                                                         AAW56763 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-JUL-1998
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                                                                                                                                                                                                                                                                                               8; Conserva
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                                                                                                                                                                                                                              DYKODDDK 8
                                                                                                                                                                                                                                                                                                                                                                    AA;
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                                                                      (first entry)
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97US-00876882.
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                                                                                                                                         peptide;
                                                                                                                                                                                                                                                                                                 100.0%; Score 47; DR
100.0%; Pred. No. 2.1
tive 0; Mismatches
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thes 0;
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     TRAIL; research; human;
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Best Local &
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                                                                                                                                          cytoplasmic :
                                                                                                                                                                                          Human LIR FLAG
             24-APR-1997;
                                                                                                         Synthetic
                                                                                                                              Immunoregulator; leukocyte immunoglobulin-like receptor; suppressor; therapeutic; treatment; disorder; IM, MMC class I receptor; inhibi cytoplasmic immunoreceptor tyrosine-based inhibitory motif; activato negative signalling; autoimmune disease; LIR.
                                                                                                                                                                                                                  09-FEB-1999
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                                 23-APR-1998;
                                                          29-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA encoding cytokine TNF-related apoptosis ligand polypeptides for producing recombinant polypeptides for research and therapy
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01-NOV-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             epitope.
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les 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                 1 DYKDDDDK 8
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                                                                                                                                                                                                                                                                                                                            DYKODDDK
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                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Wiley SR;
                                                                                                                                                                                          peptide.
            97US-00842248
                                   98WO-US008244
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95US-00548368.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORP.
                                                                                                                                                                                                                                                                                                                            8
                                                                                                                                                                                                                                                               protein;
                                                                                                                                                                                                                                                                                                                                                                                  100.0%;
                                                                                                                                                                                                                                                                8
                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                   Score 47; DB 2; Pred. No. 2.1e+06;
                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            viral infections.
                                                                                                                                                                                                                                                                                                                                                                        0,
                                                                                                                                                                                                                                                                                                                                                                                               Length 8;
                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                    inhibitor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      e apoptosis
for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               producing (TRAIL)
                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       얁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 useful
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
New nucleic acid for the antibodies and genes, use disease, specifically can
                                                                                                                                                                                  Billing-Medel PA, Co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This FLAG peptide is used in a method to isolate novel leukocyte immunoglobulin-like receptor (LIR) polypeptides. These proteins can be administered therapeutically to treat disorders associated with insufficient/defective amounts of LIR polypeptide. LIR-P3G2 and certain other LIR family members contain cytoplasmic immunoreceptor tyrosine-based inhibitory motifs (ITIMs), whilst other LIR family members lack ITIMs. By analogy with the structure and function of known MHC Class I receptor molecules, LIR's having ITIMs are inhibitory receptors mediating negative signalling, whilst those lacking ITIMs are activatory receptors. Failure of a receptor that mediates negative signalling could result in autoimmune diseases, whilst failure of a receptor mediating activatory signalling could result in suppressed immune function. They are also useful to produce probes for detecting LIR nucleic acids or isolating LIR DNA from other species
                                                                                                                                                                 Granados EN,
Russell JC,
                                                                                                           WPI; 1998-437479/37.
                                                                                                                                                                                                                                                                                                                                                                                30-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-MAR-2003
30-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                         06-AUG-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9833926-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LU105; lung disease marker;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Peptide used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW75874;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAW75874 standard; peptide; 8
                                                                                                                                                                                                                                                                          (ABBO ) ABBOTT LAB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Leukocyte immunoglobulin-like receptor, LIR, polypeptides - unifor treating autoimmune diseases or disease states associated suppressed immune function.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 18; 112pp; English.
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                                                                                                                                                           Stroupe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                   98WO-US001766
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LU105 expression
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ges SC,
ipe SD;
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                      the lung disease marker LU105 - polypeptides, used for diagnosis, prevention, treatment of lung
                                                                                                                                                                                     M, Colpitts TL, Friedman PN, Gordon o
Klass MR, Kratochvil JD, Robertsrapp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunoassay;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              lung disease; cancer; blood;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       construction
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cancer

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IT WW TUMOU
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FO 20-AL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This peptide is used in the construction of a plasmid expressing LU105 CC polypeptides. Sequences shown in AAV54616 to AAV54621 represent LU105 CC specific polynucleotide sequences which are used in the method of the CC invention for detecting target LU105 nucleic acid. The method comprises CC treating a sample with at least one LU105 specific nucleic acid, or its CC complement which is at least 50 percent identical with the LU105 specific nucleic acid sequences. LU105 is a lung disease marker. Cells transformed CC with a recombinant expression system that contains LU105 specific nucleic acid fragments, are used to express recombinant LU105 polypeptides which CC are used to raise antibodies. The antibodies are used to detect the LU105 antigen, and correspondingly this antigen is used to detect the LU105 caid sequences are used for diagnosis, staging, monitoring, prognosis, CC excid sequences are used for diagnosis, staging, monitoring, prognosis, CC glasase, specific binding agents, useful therapeutically. LU105 is a marker for lung disease (present at high concentration, in altered form CC marker for lung disease (present at high concentration, in altered form CC plasma or serum in an inexpensive, non-invasive test. (Updated on 25-MAR-CC 2003 to correct PI field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                               12-MAR-1997;
28-MAR-1997;
04-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TRAIL receptor; TRAIL-R; thrombotic thrombocytopaenic purpura; tumour necrosis factor-related apoptosis-inducing ligand; HIV infection; haemolytic uraemic syndrome; systemic lupus erythematosus; apoptosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAB99286;
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New TRAIL receptor protein and related oligomers, nucleic acid, vectors used to inhibit TRAIL activity, e.g. in cases of thrombocytic purpura, clotting in small blood vessels etc., also for diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-AUG-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-FEB-1998;
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                                                                                                                           WPI; 1998-480767/41.
                                                                                                                                                                                                                                         (IMMV ) IMMUNEX CORP
                                                                                                                                                                                                                                                                                                                                                                                                                         13-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 DYKDDDDK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DYKODDDK 8
                                                                                                                                                                                  Walczak H;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first
                                                                                                                                                                                                                                                                                              97US-00799861.
97US-00815255.
97US-00829536.
97US-00869852.
97US-00883036.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98WO-US002239.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                peptide; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               entry
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 47;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.le+06;
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      coding sequence (TRAIL-R; where TRAIL is tumour necrosis factor-related apoptosis-inducing ligand, see AAH46372 and AAB99284). TRAIL-R protein and coding sequence can be used to treat conditions involving defective or inadequate TRAIL-R, e.g. thrombotic thrombocytopaenic purpura, haemolytic-uraemic syndrome, clotting of small blood vessels, systemic lupus erythematosus and TRAIL-mediated apoptosis of T cells in human immune deficiency virus infections. The present sequence is the FLAG peptide, which was used in an example from the present invention
percent identity to the sequences shown in AAV26458 to AAV26461 are used to produce BUI01 polypeptides containing at least 1 epitope. These are used to detect BUI01-specific antibodies which are used correspondingly to detect BUI01 antigens. The BUI01 polynucleotide sequences can be used in a method for detecting the presence of a target BUI01 polynucleotide. The various assays are used for diagnosis, prognosis, staging,
                                                                                                                           The sequence encoding this peptide is incorporated into a primer for constructing a BUI01 protein expression plasmid. BUI01 is a member of the uteroglobin family of proteins and is over-expressed in breast cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to human TRAIL-receptor polypeptide and coding sequence (TRAIL-R; where TRAIL is tumour necrosis factor-relate
                                                                                               Cells transformed with a recombinant expression system comprising a sequence derived from the BU101 open reading frame and with at least 50
                                                                                                                                                                                                                                                                                                                                     Billing-Medel PA,
                                                                                                                                                                                                                                                                                                                                                                                                 19-AUG-1996;
15-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BU101; breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-MAR-2003
29-JUL-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAW54280 standard; peptide; 8 AA.
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                                                                                                                                                                                                                                       New BU101 protein over-expressed in breast cancer - useful
                                                                                                                                                                                                                                                                                                       Granados EN,
Russell JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO9807857-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Peptide sequence used in constucting a BU101 expression plasmid
                                                                                                                                                                                                                                                                      WPI; 1998-169161/15.
                                                                                                                                                                                                                                                                                                                                                                     (ABBO ) ABBOTT LAB
                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mmunisation;
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                                                                                                                                                                                           11; Page 66; 105pp; English.
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Stroupe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          grug
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97US-00912276.
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Jes SC,
Ipe SD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            screening
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                                                                                                                                                                                                                          and
                                                                                                                                                                                                                          prevention
                                                                                                                                                                                                                                                                                                                      1, Colpitts TL, Friedman PN, Gordon J
Klass MR, Kratochvil JD, Roberts-Rapp
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Pred. No.
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                                                                                                                                                                                                                            breast
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                                                                                                                                                                                                                             cancer
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RESULT 20
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Best Local
The present sequence represents a FLAG peptide for human serrate 2/ used in an example of the present invention which describes human serrate 2. The present invention also describes a method for the preparation of the polypeptides, and antibodies binding to the polypeptide and its fragments. The polypeptide and its fragments expressed by the serrate-2 gene can be used to inhibit stem (especially blood stem) cell differentiation and to inhibit endothelial cell growth. They may be incorporated in a cell culture media for culturing undifferentiated stem cells. They can also be used for treatment of neoplasms such as leukaemia. The antibodies can be used for the diagnosis of malignant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            in gene
genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           monitoring, treating and preventing diseases of the breast (especially cancer and its metastases), and also for determining susceptibility. The BUI01 polypeptides are also useful in drug screening, e.g. to identify antagonists of BUI01, potentially useful therapeutically and as targets for therapy. The antibodies are also useful for targeted drug delivery and therapeutically to neutralise BUI01 polypeptides. Fragments of the BUI01 nucleic acid are useful as probes and primers, e.g. for detection of altered gene expression or in fluorescent in situ hybridisation, also in gene therapy to generate antisense or ribozyme molecules or for
                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1998-110528/10
N-PSDB; AAV15195.
                                                                                                                                                                                                                                                                                                                         Human serrate-2 gene expression products - used to differentiation, useful in treating neoplasms, e.g.
                                                                                                                                                                                                                                                                              Example 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-JUL-1996;
14-MAY-1997;
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nia; endothelial cell; tumour; FLAG.
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97JP-00124063
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                                                                                                                                                                                                                                                                      Japanese.
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Pred. No. 2.1e+06;
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. leukaemia.
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RESULT 27
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Matches 8
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-JUN-1990;
24-AUG-1990;
13-DEC-1990;
16-MAY-1991;
12-JUL-1993;
13-MAY-1994;
                                                                                                                                II interleukin-1 receptor (IL-IN). This receptor is used in a method to investigate the regulation of the immune or inflammatory response in a mammal. This method involves administering a soluble type-II IL-1 receptor protein in an amount effective to bind to IL-1 and prevent its binding to cell-surface IL-1 receptors. The process can be used for treating alloantigen-induced rejection of transplanted tissues or organs, graft-versus-host disease and autoimmune dysfunction dependent upon the activation of T cells against self antigens, selected from rheumatoid arthritis, diabetes mellitus or multiple sclerosis
                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Regulation of interleukin-1 mediated immune or mammal - comprises administering soluble IL-1 r
                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Col
                                                                                                                                                                                                                                                                                                                                                                                                                                        treatment of e.g. graft versus
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rheumatoid arthritis;
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                                                                                                                                                                                                                                                                                                                      sequence represents an antigenic motif found in human and mouse
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8; Conser
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dysfunction; T-cell activation; self antigen;
arthritis; diabetes mellitus; multiple sclerosis.
  Conservative
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90US-00573576.

90US-00627071.

91US-00701415.

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94US-00242211.
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.0%; Score 47; Di
.0%; Pred. No. 2..
0; Mismatches
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Pred. No. 2.1e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                        d immune or inflammatory response luble IL-1 receptor protein, used disease and multiple sclerosis.
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                     2.1e+06;
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Gaps
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AAW54434
AC AAW54
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Query Match
Best Local S
Matches 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 8 AA;
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Kratochvil
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Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human PS112 expression plasmid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAW54434;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAW54434 standard; peptide;
                                                                                                                                                                                                                                                                                          AAW58571 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-APR-1998.
                                                                       Phosphatidylinositol 3-kinase; pl10 delta; human; carcinogenesis; diagnosis; PCR; primer; FLAG.
                                                                                                                                         FLAG peptide.
                                                                                                                                                                                              12-OCT-1998
                                                                                                                                                                                                                                            AAW58571;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ABBO ) ABBOTT LAB.
                      Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                рукорорк в
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               or to treat prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                    DYKDDDDDK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Friedman PN, Gordo
l JD, Roberts-Rapp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            target PS112 polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page 92; 104pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry
                                                                                                                                                                                              (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  96US-00727688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97WO-US018290
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                                                                                                                                                                                                                                                                                        peptide;
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-Rapp L, Russell i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              æ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 47; DB
Pred. No. 2.1
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              for diagnosing prostate
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Stroupe SD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
                                                                                              immune system;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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RESULT 30
AAW76424
ID AAW76
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This FLAG peptide is recognised by the M2 anti-FLAG monoclonal antibod PCR amplification of cDNA (see AAV31340) coding for human phosphatidylinositol 3-kinase (PI 3-kinase) pl10 delta subunit (see AAW38570) using a primer (see AAV31349) including a FLAG sequence yiel a FLAG-tagged PI 3-kinase construct. This was ligated into mammalian expression vector pcDNA3, and FLAG-tagged pl10 delta was expressed in transfected COS cells. pl10 delta can be used to develop products for modulating PI 3-kinase activity in immune system signalling and in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New phosphatidyl-inositol 3-kinase catalytic sub-unit - used to develop products for modulating kinase activity in immune system signalling and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chantry DH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-NOV-1997;
                                                                                                                                                                                                        JAK2 protein; cytoplasmic domain; betac subunit; screening; ast interleukin; granulocyte macrophage-colony stimulating factor; IL-3; IL-5; human; FLAG.
                                                                                                                                                                                                                                                                                                 AAW76424;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-322736/28
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                                                                                                                                           01-OCT-1998
                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                       FLAG affinity
                                                                                                                                                                                                                                                                           20-JAN-1999
                                                                                                                                                                                                                                                                                                                      AAW76424 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         in carcinogenesis.
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                               WPI; 1998-532151/45
                                                                                               24-MAR-1997;
                                                                                                                     23-MAR-1998;
                                                                                                                                                                WO9843087-A1
                                                                         (PHAA)
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                                                                         PHARMACIA & UPJOHN CO
                                                                                                                                                                                                                                                                                                                                                                              DYKDDDDK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 12; 53pp;
                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
ilarity 100.0%;
Conservative C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hoekstra
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                                                   Kaytes
                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                       label used in the course
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                                                                                               97US-0041511P
                                                                                                                     98WO-US005387.
                                                                                                                                                                                                                                                                                                                      peptide; 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Holtzman
                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                        Score 47; Db 2, -
Pred. No. 2.1e+06;
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                                                                                                                                                                                                                                                        of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 8;
                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                         GM-CSF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               yielded
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Screening for compounds useful for preventing or treating asthma -determining if compounds inhibit binding of the JAK2 protein to e.s

e.g.

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GM-CSF

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RESULT 31
AAW81752
ID AAW81752
ID AC AAW81
AC AA
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Best Local
                                              This sequence is a FLAG peptide used in a method resulting in the isolation of a US6 protein from human cytomegalovirus which can be used as therapeutic or prophylactic immunomodulators, in particular inhibitors of antigen presentation by MHC Class I on the surface of cells. A particularly application of this protein is to reduce immunogenicity of cells being transfected for gene therapy. Fragments of US6 nucleic acid and is analogues are useful diagnostically as probes and primers, and also as antisense sequences and ribozymes. US6 proteins are used to raise antibodies. The US6 protein inhibits a later stage of antigen
                                presentation than
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New US6 gene of cytomegalovirus - useful for diagnosis and therapeutic prophylactic immunomodulation, e.g. to reduce immunogenicity of gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US6; immunomodulator; therapeutic; prophylactic; inhibitor; cell surface; antigen presentation; major histocompatibility complex; MHC Class I; immunogenicity; gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This represents a FLAG affinity label contained in a recombinant JAK2 protein used in the course of the invention. The invention provides a method of screening for compounds useful for treating or preventing asthma. The method comprises contacting a molecule comprising at least the N-terminal 294 amino acid residues of the JAK2 protein, with another molecule comprising at least 13 membrane-proximal cytoplasmic amino acids of interleukin (II)-3, IL-5 or granulocyte macrophage-colony stimulating factor (GM-CSF) proteins in the presence of the candidate compound, and determining whether the first and the second molecules form a complex. If the compound inhibits complex formation, it can be used to treat asthma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1998-583601/49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (BOEF ) BOEHRINGER MANNHEIM GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-JAN-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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                                known inhibitors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein; 8
                                                                                                                                                                                                                                                                                                                                                                                                       34pp; German
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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RESULT 32
AAW79682
ID AAW79
XX AAW79
XX AAW79
XX III-JA
XX Gastr
KW Gastr
KW Human
XX Homo
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Matches
                                                                                           Matches
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                                                                                                                                                                                                                                                   This sequence represents a C-terminal fragment of the human CS198 protein which is used in a method to detect the presence of a target CS198 polynucleotide in a test sample. The CS198 gene is useful as a marker for gastrointestinal (GI) tract disorders. The methods and products can be used in detecting, diagnosing, staging, monitoring, prognosticating, preventing or treating, or determining the predisposition to diseases and conditions of the GI tract, such as GI tract cancer, Barret's ossophagus, gastric ulcer, gastritis, leiomyoma, polyps, Crohn's disease, ulcerative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New gastrointestinal polynucleotides, CS198, and their detection - used for developing products for the diagnosis and treatment of gastrointestinal disorders, e.g. cancers, gastric ulcer or gastritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Billing-Medel PA,
Granados EN, Hay
Roberts-Rapp L,
                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                       gastric ulcer, colitis, and pa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 11a; Page 104; 127pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gastrointestinal tract; GI tract; cancer; disease; detection; human; predisposition; treatment; Barret's oesophagus; gastric gastritis; lelomyoma; polyps; Crohn's disease; ulcerative col:
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8; Conserv
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                                                                                              Conservative
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L, Russell
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                                                                                                             100.0%;
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JC, Stroupe
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lges SC, K
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Pred. No.
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Pred. No.
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Klass
pe SD;
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hes 0;
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MR, Kratoch
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                                                                                                                                        Length 8;
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Kratochvil JD;
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RESULT 33 AAW79770

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Matches
                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a C-terminal fragment from the human 20 kD presenilin 2 protein (PS2-CTF) which is useful in screening for compounds that inhibit proteolytic processing of PS2 in cells. Tracking of presenilin 2 or PS2-CTF, when expressed in cells and in screening assays, is done by linking a FLAG epitope peptide (this sequence) to the N- or C-terminus of the PS2 protein. The inhibitory compounds can be administered to prevent proteolytic cleavage of presenilin 2 at a cleavage site which generates PS2-CTF, and therefore to inhibit apoptotic cell death. Mutations in the presenilin genes presenilin 1 and presenilin 2 (PS1 and PS2) cause a major portion (at least 50%) of early onset familial Alzheimer's disease (FAD), and detection of abnormal PS2-CTF leevels can be indicative of a specific disease, e.g. FAD. (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (TANZ/)
(KIMT/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20 kDa presenilin 2 C-terminal fragment, PS2-CTF - useful to screen compounds inhibiting proteolysis of presenilin 2 and to produce antibodies used in diagnosis of familial Alzheimer's.
                                                                                                                                                                                                                                                                                                                                               Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tanzi RE,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Proteolytic processing; human; presenilin 2; inhibitor; apoptosis; cell death; early onset familial Alzheimer's disease; FAD; diagnosis.
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02-MAR-1999
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Chimeric; adenovirus; fiber protein; binding; targeting; coat protein; constrained peptide motif; gene therapy; cancer; heart disease; autoimmune disorder.
                                                           Chimeric adenovirus fiber protein non-native amino acid sequence
                                                                                        29-JUL-1998
                                                                                                                                                AAW56033 standard; peptide; 8
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KIM T.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 38; 84pp;
                                                                                                                                                                                                                                                                                   Conservative
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Pred. No.
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Wickham TJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-AUG-1997;
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Mastadenovirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 7; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence
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                       28-MAR-1997;
09-MAY-1997;
                                                                                                                                                             ZSIG-35; beta-chemokine; human; ligand; lymphocyte migration; inflammation; ischaemia; reperfusion injury; FLAG.
                                                                                                                                                                                                                            02-FEB-1999
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                                                                                                                                     Synthetic.
                                                                                                                                                                                                   FLAG tag
                                                                                                                                                                                                                                                     AAW30566;
                                                                                                                                                                                                                                                                              AAW30566 standard;
                                                                                                              WO9844117-A1
(ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                                                                                                            1 DYKDDDDK 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         adenovirus fibre proteins - containing non-native amino acid co provide for binding and entry into cells, especially for gene
                                                                                                                                                                                                                                                                                                                                             DYKODDDK
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                       97US-0042862P.
97US-0046083P.
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                                                                                                                                                                                                                                                                              peptide; 8 AA.
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hes 0;
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RESULT 36
AAM75806
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                                                                                                                                                                                                                                                                           Tryptase-6 complex inhibitory peptides - used to treat mast cell-mediated inflammatory disorders e.g. asthma.
                       This represents a FLAG peptide epitope sequence. The invention provides sequences shown in AAW63160 to AAW63169 that are inhibitors of mouse mast cell protease (mMCP-6). These peptides which are tryptase-6 complex inhibitors, can be used for treating a mast cell-mediated inflammatory disorder. The inhibitors can be used to treat inflammatory disorders including asthma, allergic rhinitis, urticaria and antioedema, eczematous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mast cell protease; MCP; mouse; inhibitor; peptide substrate; asthma; tryptase-6 protein; inflammatory disorder; allergic rhinitis; urticaria; antioedema; eczematous dermatitis; atopic dermatitis; anaphylaxis; hyperproliferative skin disease; peptic ulcer; hyperresponsiveness; inflammatory skin condition; FLAG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This FLAG peptide can be used as a N- or C-terminal tag of novel human chemokine 2SIG-35 polypeptides (see AAW30565) of the invention. Vectors have been designed for expression of FLAG-tagged ZSIG-35 in mammalian host cells. The FLAG peptide acts as a purification aid. ZSIG-35 polypeptides are useful in the regulation of acute and chronic inflammatory disease conditions, lymphocyte migration and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-FEB-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BRIGHAM & WOMENS HOSPITAL
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                                                                                                                                                                                                                      Page 18; 69pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard;
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  (atopic
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                                                                                                                                                                                                                                                                                                                                                                                                                   Huang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     epitope
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              97US-0037090P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98WO-US001865
  dermatitis),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    peptide; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZSIG-35 - used for, e.g. treating inflammatory migration and ischaemia/reperfusion injury.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          8
anaphylaxis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.1e+06;
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hyperproliferative
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                                           Query Match
Best Local S
Matches 8
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Best Local
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                                                                                                                       mutations that qualitatively alter the protein product or any protein encoding gene, using genomic DNA or cDNA as the starting material. The method is useful for identification of mutations in certain disease
                                                                                                                                                      The DNA sequences (AAV26452-V26455) are examples of 4 elements which present in a unique oligonucleotide sequence. The Flag peptide is an example of a tag which can be encoded by a region of the unique oligonucleotide sequence. The unique oligonucleotide sequence is used amplify desired mutated sequences and to tag the encoded peptide at terminus. Ligands recognising the peptide can be used to purify the proteins for further analysis. The method is capable of detecting
                                                                                       Sequence 8
                                                                                                                                                                                                                                                                                   Mutation detection using peptide tagged in vitro synthesised proteins useful for identification of mutations in disease causing genes, e.g.
                                                                                                              causing
                                                                                                                                                                                                                                                  Disclosure; Page 3;
                                                                                                                                                                                                                                                                        breast cancer susceptibility genes
                                                                                                                                                                                                                                                                                                                                  WPI; 1998-207407/18
                                                                                                                                                                                                                                                                                                                                                         Garvin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FLAG
                                                                                                                                                                                                                                                                                                                                                                                                     13-SEP-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                               (GARV/) GARVIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9811249-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Peptide tag; | BRCA1; BRCA2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAW54266
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAW54266 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 disease, peptic ulcers, inflammatory bowel disorder, hyperresponsiveness and inflammatory skin conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 DYKDDDDK 8
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8; Conserv
                                            8
                                                                                                              genes,
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                                                       Similarity
DYKDDDDDK
                      DYKDDDDK
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                                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               purification; identification;
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                                                                                                              e.g.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amplification; primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        peptide; 8- AA.
                                                                                                                                                                                                                                                  13pp; English.
                                                                                                            the breast cancer susceptibility genes
                                                     100.0%;
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                                           Score 47; DE
Pred. No. 2.1
0; Mismatches
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                                                                                                                                                                                                                                                                         BRCA1
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No.
                                                     2.1e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mutation; breast cancer;
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RESULT 39
AAW79562
ID AAW79
XX
AC AAW79
XX
DT 11-JA
XX
DE FLAG
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                                                                                                                                                                                                                                                                                                                            The synthetic octapeptide recognised by anti-FLAG M2 was used in the CC creation of a plasmid for the expression of secretable TNF-related CC endothelium proliferative agent (TREPA). TREPA, or its activators or CC agonists, are used to treat a deficit of TREPA, e.g. to promote wound CC healing or tissue grafting, by promoting vascularisation, also to induce CC apoptosis for treating cancer and eliminating autoreactive T cells, as an CC adjunct to cancer chemotherapy or antiviral treatment. TREPA peptides can CC also be used to target cytotoxic agents or for affinity isolation of the CC corresponding receptor, the nucleic acid for which can be used to treat TREPA minics. Ribozymes, antisense RNA, antibodies or peptides, are used to treat TREPA-associated diseases, e.g. tumours and CC metastases (by inhibiting vascularisation), inflammation or a wide range of autoimmune conditions, conditions involving abnormal stimulation of CC cyulation and placental formation) or other angiogenic conditions (e.g.
                                                                                                                                                                                                                            Query Match
Best Local S
Matches 8
                                                                                                                                                                                                                                                                                        Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 43; Page 126; 142pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Detecting nucleic acid encoding TREPA - useful for diagnosis and treatment of autoimmune disease, tumours and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1998-447255/38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-FEB-1997;
10-FEB-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TNF; endothelium proliferative agent; TREPA; wound healing; cancer; tissue grafting; vascularisation; apoptosis; autoimmune; birth control.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic octapeptide recognised by anti-FLAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-OCT-1998
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                             11-JAN-1999
FLAG peptide
                                                             AAW79562;
                                                                                       AAW79562 standard; peptide; 8
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                                                                                                                                                                                   1 DYKDDDDK 8
                                                                                                                                                                                                                            8;
                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                  DYKDDDDK 8
                                                                                                                                                                                                                            Conservative
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                             (first entry)
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98US-00021706.
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                                                                                                                                                                                                                                         100.0%; Score 47;
100.0%; Pred. No.
                                                                                                                                                                                                                            <u>.</u>
                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                          DB 2;
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                                                                                                                                                                                                                                                        Length 8;
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22-DEC-1997;

97WO-US023775

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RESULT 40
AAW69953
ID AAW69
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This is the amino acid sequence of a FLAG peptide. DNA encoding FLAG (see AAV61524) can be introduced into an adenovirus vector. The invention provides a chimeric adenovirus locat protein, particularly a chimeric adenovirus hexon protein (see AAW79538-39), that has a decreased ability or inability to be recognised by a neutralising antibody directed against the corresponding wild-type adenovirus coat protein. The chimeric adenovirus coat protein has a non-native amino acid sequence, especially comprising a deletion of one or more regions of the hexon protein. A vector that comprises the chimeric adenovirus coat protein is claimed,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chimeric adenovirus coat protein - useful in, transfer to treat inherited genetic diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                              RANK; necrosis factor-kappa B; NF-kB; receptor activator; human; immune response; inflammatory response; toxic shock; sepsis; TNF RANK ligand; tumour necrosis factor; FLAG peptide epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 8 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-MAR-1997;
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                                                                                                                                                                                                                                                                                                           08-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                     AAW69953 standard; peptide; 8 AA
                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                          FLAG peptide
                                                                                                                                                                                                                                                                                                                                                         AAW69953;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and can be used for gene transfer, for the treatment of inherited
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CORR ) CORNELL RES FOUND INC (GENV-) GENVEC INC.
02-JUL-1998
                                              WO9828426-A2
                                                                                          Tomo sapiens
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                                                                                                                                                                                                                                                          for constructing RANK/Fc
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Pred. No. 2.1e+06;
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                                                                                                                                                                                                                                                               fusion
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RESULT 41
AAW48629
ID AAW48
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CC This represents a FLAG peptide epitope used in the construction of a CC RANK/FC fusion protein. RANK (receptor activator of necrosis factor-CC kappaB (NF-KB)) is a member of the tumour necrosis factor (TNF) family. A CC soluble form of RANK is fused to the FC region of human IgGl. A soluble CC RANK may be used for inhibiting activation of NF-KB, by contacting a cell expressing membrane-associated RANK with a soluble RANK which binds to CC RANK ligand (RANKL). RANKL polypeptides can activate RANK which binds to CC to induce maturation of dendritic cells and enhance their allocation in the composition may also be used for regulating an immune or CC inflammatory response. Inhibition of NF-KB prank tragonists may be cuseful in ameliorating negative effects of an inflammatory response that result from triggering of RANK, e.g. in treating toxic shock or sepsis, CC graft-versus-host reactions, or acute inflammatory reactions. They can cells that express RANK. RANKL polypeptides can also be used to identify cinhibitors of RANK and thus inhibitors of an inflammatory response, and CC class of reprotecting RANK-expressing cells from the negative effects of CC chemotherapy or the presence of high levels of TNF-alpha. The products of CC can also be used for detection and drug screening
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Best Local S
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07-MAR-1997;
14-OCT-1997;
                                                                                                                                                                                                                                                Chimeric adenoviral penton base protein epitope
                                                                                                                                         WO9807877-A1
                                                                                                                                                                        Synthetic
                                                                                                                                                                                                        bispecific
                                                                                                                                                                                                                  Short-shafted adenoviral fibre; ssAF; penton base; cystic fibrosis;
                                                                                                                                                                                                                                                                                       11-SEP-1998
                                                                                                                                                                                                                                                                                                                                                 AAW48629 standard; peptide; 8 AA
               (GENV-) GENVEC INC
                                           21-AUG-1996;
                                                                         21-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New isolated ligand for receptor activator of NF-kappa B - used to develop products for augmenting an immune response for inhibiting an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 6; Page 53; 80pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Anderson DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        develop products for augmenting an immune response inflammatory response and for protection of cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1998-377657/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (IMMV ) IMMUNEX CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
les 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 рукрарак в
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                    рукрарак в
                                                                                                                                                                                                    antibody; HIV; cancer
                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Galibert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    96US-0059978P.
97US-00813509.
97US-0064671P.
                                           96US-00700846
                                                                         97WO-US014718
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ٥,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Maraskovsky
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 47; DB z; Pred. No. 2.1e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
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RESULT 42 AAW47353
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence is a FLAG octapeptide which is incorporated into an CC adenoviral fiber. The fiber protein and its penton base are used by the cadenovirus to interact with distinct cellular receptors to attach to and CC efficiently infect a cell. The invention claims for a transfer vector CC which contains DNA encoding a short-shafted adenoviral fibre (ssAF). The cadenoviral penton base can be modified by incorporating sequences for a cligand to a cell surface receptor or sequences that allow binding to a cell surface receptor and the sequence and the other end to bind specific antibody molecule to bind to the fiber can cotapeptide allows one end of the antibody molecule to bind to the fiber can the other end to bind specifically to a cell surface receptor. Therefore this method enables targeting of the adenovirus to a desired CC cell-surface receptor by the introduction of a non-native amino acid sequence can be such that it enables direct or indirect binding, e.g. by means of a bispecific or multispecific binding agent, of the adenovirus to the pentor base or the fibre knob. The non-native amino acid sequence can be such that it enables direct or indirect binding agent, of the adenovirus to the sactive of a bispecific or multispecific binding agent, of the adenovirus to the sactive of a bispecific or multispecific binding agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local (
                                                                                                                                                                                                                                                                             EBV; BZLF2; beta-chain; major histocompatibility complex; MHC; class II; antigen; prevention; treatment; autoimmune disease; transplant rejection allergy; asthma; super-antigen; Epstein-barr virus; FLAG-peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the adenovirus to the desired cell-surface receptor. Using the a short shafted adenovirus to attach to a cell and therefore effect entry into the cell would reduce the level or efficiency of adenoviral fibre binding to its cell-surface receptor and increase the adenoviral penton base binding to its cell-surface receptor, thereby increasing the specificity of binding of the adenovirus to a given cell. The transfer vector is claimed to be useful for expressing therapeutic genes, for treating inherited diseases such as cystic fibrosis or for inducing cell death in pathogenic infections (e.g. HIV infections) and cancers
                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                              FLAG-peptide.
                                                                                                                                                                                                                                                                                                                                                                                                         01-JUN-1998
                                                                          28-APR-1994;
                                                                                                                   28-APR-1995;
                                                                                                                                                          10-MAR-1998
                                                                                                                                                                                                US5726286-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW47353 standard; peptide; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 8 AA;
                                    (IMMV ) IMMUNEX CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW47353;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Short-shafted adenoviral fibre constructs - used for producing products that facilitate entry into target cells, useful for gene therapy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cellular disorders.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wickham TJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 рукорорк в
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DYKODDDK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Roelvink PW,
                                                                        94US-00235397
                                                                                                                   95US-00430633
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kovesdi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.1e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 2;
                                                                                                                                                                                                                                                                           disease; transplant re
rr virus; FLAG-peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 8;
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                                                                                                                                                                                                                                                                                                  rejection,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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Comeau MR,

Alderson M,

Farrah TM,

Hutt-Fletcher LM,

Armitage

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ARBSULT 43
AAW76407
ID AAW76
XX AAW76
AC AAW76
AC AAW76
XX DT 07-DE
XX PAR-2
KW treat
KW treat
KW treat
KW intes
XX Synth
PN WO984
PN WO984
PF 27-MP
XX PREG
XX PREG
XX PREG
XX Preve
PT Prote
XX Preve
PT Prote
XX This
PS Examm
XX This
CC epitc
CC operx
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PAR-2; proteinase-activated treatment; prevention; mast chymase; Ca2+ mobilisation; intestinal motility; colon;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence was used in the preparation of a synthetic construct containing the Epstein-barr virus (EBV) BZLF2 protein, which is capable of binding a beta-chain of a major histocompatibility complex (MRC) class II antigen. The protein can be used to inhibit antigen-specific antibody formation, peripheral blood mononuclear cell proliferation and cytotoxic T-cell responses, e.g. in the prevention or treatment of autoimmune diseases, transplant rejection, allergies or asthma, and as a super-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Col 31-32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 transplant rejection.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Spriggs MK,
                                                                                                                                                                                                                                                                                                                                                                                                              Bunnett NW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9843477-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human PAR-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW76407;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAW76407 standard;
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prevents mast cell degranulation, inhibits tryptase/chymase activity or antagonises PAR-2. PAR-2 fragments are also used in a method for identifying candidate compounds for use in treating POI, by screening for the compounds for use in treating POI, by screening for the compounds for use in treating POI, by screening for the compounds for use in treating POI, by screening for the compounds for use in treating POI, by screening for the compounds for use in treating POI, by screening for the compounds for use in treating POI, by screening for use in the point POI in the p
                                                                      This sequence is a human proteinase-activated receptor, PAR-2, epitope which is used in a method for preventing or treating poperative ileus (POI) in a mammal by administering a compound
                                                                                                                                                                                                                                                                    Preventing or treating surgically induced ileus - prevent mast cell degeneration, inhibit tryptase o
                                                                                                                                                                                                                                                proteinase-activated receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (REGC ) UNIV
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                                                                                                                                                                                     6; Page 12; 20pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
8; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                              Corvera CU
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 peptide; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        receptor 2; post-operative ilius; POI; cell degranulation; inhibitor; tryptastrypsin; activating peptide; surgery; Flag epitope.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                         tryptase or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.1e+06;
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                                                                                                                                                                                                                                                                       using compounds which
or chymase or antagonise
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                                                                                                     post-
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                                                                                                                                    Flag
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RESULT 44
AAW56251
ID AAW56
XX AAW56
AC AAW56
XX Thera
XX Thera
XX Thera
XX Thera
XX Synth
XX Synth
XX II-SE
PR 10-SE
PR 10-SE
PR 27-FE
XX II-SE
PR 27-FE
XX II-SE
PR 27-FE
XX II-SE
PR 10-SE
PR
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Best Local :
                                                                                 The FLAG peptide was used to tag recombinant murine Interleukine-13 (II-13) in the production of II-13 binding protein, and related, therapeutic molecules. The therapeutic molecules can be used in the antagonism of at least one II-13 activity. They can be used for treating II-13 mediated conditions such as certain allergic conditions such as asthma or to inactivate locally administered II-13 after II-13 treatment. The products can also be used as diagnostic agents, e.g. for detecting autoimmune diseases. The antibodies can also be used for immunotherapy and may also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the ability to inhibit Ca2+ mobilisation in cells transfected with the PAR-2 gene when the cells are stimulated by trypsin or activating peptide. A compound is selected when there is significant inhibition of Ca2+ mobilisation in the cells when compared with activated cells in the absence of the compound. This method can be used to prevent or treat inhibition of intestinal (especially colonic) motility which can result inhibition of intestinal (especially colonic) motility which can result
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     from abdominal surgery. Unlike current treatments, this method is effective, non-invasive and avoids the need for secondary surgical intervention and prolonged hospitalisation. In addition, recovery time
   Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated interleukin-13 binding protein for therapy e.g. for allergic conditions suc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1998-207062/18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nicola
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27-FEB-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Therapeutic; IL-3 mediated condition; allergy; asthma; diagnosis; autoimmune disease; antibody; immunotherapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAW56251;
                                                                                                                                                                                                                                                                                                                                              Example 1; Page 28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (AMRA-) AMRAD OPERATIONS PTY LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Euthoritos
                                                            used
                                                                                                                                                                                                                                                                                                                                                                                                                detection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    8
                                                         as a diagnostic tool
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DYKDDDDK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DYKODDDDK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                colcnic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hilton DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     96AU-00002262.
97AU-00005374.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             surgery is accelerated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein;
                                                                                                                                                                                                                                                                                                                                                 69pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Zhang J,
                                                                                                                                                                                                                                                                                                                                                    English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              œ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 47; DB Pred. No. 2.1 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Simpson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; DB 2;
2.1e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                        ein -
such
                                                                                                                                                                                                                                                                                                                                                                                                                                        used to develop products as asthma or for diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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Query Match Best Local S Matches 8

Similarity 8; Conser

Conservative

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100.0%;

Score 47; DB Pred. No. 2.1 0; Mismatches

2.1e+06;

Indels 8

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Gaps

0

DB 2;

Length

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UXAX EXEXEX EXXXXX
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AAW46969
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                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 8
                                 RANK; necrosis factor-kappa B; NF-kB; receptor activator; human; immune response; inflammatory response; toxic shock; sepsis; TNF; RANKL; RANK ligand; tumour necrosis factor; FLAG peptide epitope.
                                                                                                                                                                                                                                                                                                                   describes a novel human slit-like protein (the mature protein in Claim 1). The slit-like polypeptide is useful for diagnosis treatment of brain-specific diseases and cancers. Antibodies diagnist the protein, or its fragments can also be used for diagnosts.
                    Synthetic.
                                                                              FLAG peptide epitope for constructing
                                                                                                     08-OCT-1998
                                                                                                                                            AAW68289 standard; peptide;
                                                                                                                                                                                                                                                                                      Sequence 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Slit-like protein; human; diagnosis; treatment; brain-specific disease; cancer; antibody.
       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                   Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                               Human Slit-like protein - useful for diagnosis and treatment of brain-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-APR-1998.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Peptide sequence of the specification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-JUL-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW46969 standard; protein; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ASAH ) ASAHI KASEI KOGYO KK
                                                                                                                                                                                                                  1 DYKODDDK 8
                                                                                                                                                                                                                                          8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ш
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                                                                                                                                                                                                   DYKDDDDK 8
                                                                                                                                                                                                                                                                                                                                                                                                       diseases and
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                                                                                                                                                                                                                                                                                     A,
                                                                                                                                                                                                                                          100.0%;
ilarity 100.0%;
Conservative 0
                                                                                                                                                                                                                                                                                                                                                              sequence
                                                                                                                                                                                                                                                                                                                                                                                   Page 39;
                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 96JP-00186219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     97JP-00205351
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                                                                                                                                                                                                                                                                                                                                                              appears in the specification.
                                                                                                                                                                                                                                                                                                                                                                                 45pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                        cancers.
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AA
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Pred. No.
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                                                                            RANK/Fc fusion protein.
                                                                                                                                                                                                                                          2.1e+06;
thes 0;
                                                                                                                                                                                                                                                               DB 2;
                                                                                                                                                                                                                                                              Length 8;
                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                          The specification
                                                                                                                                                                                                                                          0,
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                                                                                                                                                                                                                                         Gaps
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AAW44008
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Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Host cells transformed or transfected with an expression vector comprising the RANK encoding nucleic acid can be used to produce recombinant RANK protein. A soluble form of RANK is fused to the FC region of human IgG1. The soluble farm of RANK is fused for inhibiting activation of NF-kB, by contacting a cell expressing membrane-associated RANK with a soluble RANK which binds to RANK ligand (RANKL). The soluble RANK polypeptide composition may also be used for regulating an immune or inflammatory response. Inhibition of NF-kB by RANK antagonists may be useful in ameliorating negative effects of an inflammatory response that result from triggering of RANK, e.g. in treating toxic shock or sepsis, graft-versus-host reactions, or acute inflammatory reactions. They can also be used in adjunct therapy for disease characterised by neoplastic cells that express RANK. The products can also be used for detection and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-DEC-1996;
07-MAR-1997;
14-OCT-1997;
                                                                                                                                         Prion; epitope; FLAG; Strep; poly-histidine; transgenic animal; scrapie; Creutzfeldt-Jakob
                                         11-DEC-1997.
                                                                                                Synthetic
                                                                                                                                                                                 FLAG peptide I used in an epitope tagged prion protein construct
                                                                                                                                                                                                                   05-JUN-1998
                                                                                                                                                                                                                                                                        AAW44008 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 8 AA;
                                                                                                                             bovine
                                                                                                                                                                                                                                                AAW44008;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  drug screening
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New isolated receptor activator of necrosis factor-kappa B - usefie.g. developing products for regulating an immune or inflammatory response, treating toxic shock or sepsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This represents a FLAG peptide epitope used in the construction of a RANK/FC fusion protein. RANK (receptor activator of necrosis factor-kappaB (NF-kB)) is a member of the tumour necrosis factor (TNF) family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 6; Page 53; 80pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Anderson DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-DEC-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (IMMV ) IMMUNEX CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1998-377655/32
                                                                                                                           spongiform encephalopathy; BSE.
                                                                                                                                                                                                                                                                                                                                                                               1 рукророк в
                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
8; Conserv
                                                                                                                                                                                                                                                                                                                                                   DYKDDDDK
                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                   (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Galibert
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97US-00813509.
97US-0064671P.
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                                                                                                                                                                                                                                                                                                                                                   8
                                                                                                                                                                                                                 entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                          .0%; Score 47;
                                                                                                                                                                                                                                                                           ω
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                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                          . 2.1e+06;
ches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    M
                                                                                                                                                                                                                                                                                                                                                                                                                                         B
                                                                                                                                           haemagglutinin;
b disease; CJD;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 8;
                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           B - useful for,
                                                                                                                                                                                                                                                                                                                                                                                                          0,
                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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29-MAY-1997;

97WO-US009289

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RESULT 48
AAW80478
ID AAW80
XX AAW80
XX AAW80
XX AAW80
XX Pepti
XX EST s
XX Lymph
XX Synth
XX AAW80
PN WO984
XX Synth
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CC encoding an amino acid sequence of a biologically active protein fragment CC and a second nucleic acid sequence encoding a heterologous epitope domain is a peptide selected from a CC domain. The heterologous epitope domain is a peptide selected from a CC peptide group of FLAG, Strep, poly-histidine, human c-myc peptide cc recognised by monoclonal antibody 9E10 and haemagglutinin peptide cc recognised by monoclonal antibody 12CAS. The protein has two different three cc recognised by monoclonal antibody 12CAS. The protein has two different three cc synthetic or chimeric PrP molecule. The protein has two different three cc antimation relative to the protein such that the epitope domain is spatially positioned cc relative to the protein such that the epitope domain is more exposed in a first confirmation relative to a second conformation. The nucleic acid construct may be used for the production of transgenic animals or cells conformation and shapes of a protein. These methods are particularly useful in studying diseases caused by prion proteins, e.g. Creutzfeldt-Jakob cc disease (CJD), scrapie and bovine spongiform encephalopathy (BSE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAW80478;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW80478 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This sequence represents an artificial FLAG peptide epitope. It is us in a recombinant nucleic acid construct encoding an epitope-tagged poptotein (PrP). The construct comprises a first nucleic acid sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     especially epitope-tagged
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nucleic
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  WPI; 1998-557139/47
                                                                        Billing-Medel PA,
Granados EN, Hayı
                                                                                                                                                                                                  31-MAR-1997;
                                                                                                                                                                                                                                                 31-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                    WO9844160-A1
                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Peptide sequence encoded by an antisense primer.
                                              Roberts-Rapp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EST sequence;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lymphoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1998-042112/04
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8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              VIND
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                                                                                                                                                    ABBOTT
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nilarity 100.
Conservative
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                                                                      Hayden M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CS110;
                                                                                                                                                    LAB.
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                                                                                                                                                                                                  97US-00829755
                                                                                                                                                                                                                                                    98WO-US006338
                                                 Russell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            peptide; 8
                                                                                             Cohen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gastrointestinal tract;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            с
С,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                . 0%
                                            M, Colpitts TL, Hodges SC, Klass JC, Stroupe SD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  prion
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein
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                                                                                             TL,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .le+06;
                                                                        Friedman PN,
MR, Kratoch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ¥
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cancer; adenocarcinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                           Kratochvil JD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                     Gordon J;
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RESULT 49
AAW70592
ID AAW70
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Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence represents a peptide sequence encoded by an antisense primer of the invention. The sequence incorporates a recognition site to aid in analysis and purification of the CS110 protein. A set of contiguous and partially overlapping EST sequences (AAV63521-27) are designated CS110. The sequences are isolated from a CDNA library made from gastrointestinal tract tumour and normal tissues. The polypeptides are useful diagnostically to detect CS110 antigen/anticCS110 antibody in samples, e.g. to detect diseases and conditions of the gastrointestinal tract, especially cancers, e.g. adenocarcinoma and
The present sequence represents a Flag sequence used in the course of t invention. The specification describes zsig25 protein, an adhesion-modulating protein expressed at very high level in prostate cells. The protein is useful as a diagnostic marker for prostatic carcinoma and B-cell cancers, possibly also for infertility, and as a reagent for separating cancerous and non-cancerous cells. The products may also be used to diagnose or treat Wolf-Hirschhorn syndrome, associated with a deletion in the region of chromosome 4 (p16.3) where the zsig25 gene is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       zsig25; adhesion-modulating
B-cell cancer; infertility;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAW70592 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New CS110 polypeptide(s) - useful for diseases, e.g. gastrointestinal tract
                                                                                                                                                                                      used
                                                                                                                                                                                                                                                                                                                      10-APR-1997;
11-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                             WO9845442-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     chromosome 4 (p16.3).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Flag sequence used in the course of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW70592;
                                                                                                                                         Claim
                                                                                                                                                                   stimulation of haematopoietic cells,
                                                                                                                                                                                                 New zsig25
                                                                                                                                                                                                                                WPI; 1998-557522/47.
                                                                                                                                                                                                                                                         Sheppard
                                                                                                                                                                                                                                                                                                                                                                                                15-OCT-1998
                                                                                                                                                                                                                                                                                         (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                                                                                                                   10-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 lymphoma
                                                                                                                                                                     zsig25 protein - and antibodies, involved in modulation of adhesion, for diagnosis and treatment of prostatic and B-cell tumours, ulation of haematopoietic cells, treatment of immune deficiency etc.
                                                                                                                                         10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 рукрарак в
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8; Conserv
                                                                                                                                                                                                                                                                Ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DYKDDDDK
                                                                                                                                       Page 120; 161pp; English.
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97US-0049288P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        peptide;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     protein; prostate cell; prostatic carcinoma;
Wolf-Hirschhorn syndrome;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 47; DB Pred. No. 2.1 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ng gastrointestinal tract and to produce antibodies.
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                                                                                                                course of the
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RESULT 50
AAE08734
ID AAE08
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       The patent discloses novel receptor activator of nuclear factor (NF)-
kappab (RANK) proteins and their corresponding DNAs. RANK is a member of
the tumour necrosis factor (TNF) receptor superfamily and associates with
TNF receptor associated factor (TRAF) 2 and 3 which are important in the
regulation of immune and inflammatory response. The receptors are useful
for regulating immune response and in screening for inhibitors of these
receptors. The cytoplasmic domain of RANK is used in developing assays
for inhibit interaction of RANK with TRAF1, TRAF2, TRAF3, TRAF5 and
particularly TRAF6. NF-kappab inhibition by RANK antagonists are useful
in ameliorating the negative effects of an inflammatory response that
result from triggering of RANK, e.g. in treating toxic shock or sepsis,
graft-versus-host reactions, acute inflammatory reactions and the effects
of bone resorption. RANK acts as an anti- apoptotic signal and rescue the
cells that express RANK from apoptosis. Soluble forms of the receptor are
used in vivo or in vitro based screening tests for agonists or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-DEC-1996;
23-DEC-1996;
07-MAR-1997;
                                                                                                                                                                                                                                                                                                     New isolated receptor activator of necrosis factor-kappa B - useful e.g. developing products for regulating an immune or inflammatory response, treating toxic shock or sepsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; receptor activator of nuclear factor kappaB; RANK; NF; tumour necrosis factor; TNF; TNF receptor associated factor; TRAF; immune response; inflammatory response; graft-versus-host reaction; toxic shock; sepsis; acute inflammatory reaction; bone resorption;
                                                                                                                                                                                                                                                                             Example 6; Col
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-377655/32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-MAR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FLAG tag, used to determine the association of TRAF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-NOV-2001
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96US-00772330.
97US-0077181P.
97US-00813509.
97US-0064671P.
97US-00996139.
                                                                                                                                                                                                                                                                         59; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             signal; therapy; immunosuppressant; anti-inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98US-00215649
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Pred. No. 2.1e+06;
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 antagonists
 of RANK-mediated
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NF-kappa
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RESULT 51
AAW94672
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    activation, or to inhibit transduction of a signal via RANK. RANK compositions are used in the development of both agonistic and antagonistic antibodies, or as an adjunct therapy for disease characterised by neoplastic cells that express RANK. Compounds that interfere with RANK/TRAF6 interactions are useful for modulating the formation of osteoclasts from osteoclast precursors and for modulating osteoclast function and activities. They are useful as inhibitors of diseases associated with excess bone resorption and as immunosuppressants or anti-inflammatory agents. The RANK DNAs are useful for the expression of recombinant proteins, as probes for analysis of the presence or distribution of RANK transcripts, while the proteins are useful in preparing kits for the detection of soluble RANK, or monitor RANK-related activity. The present sequence is a FLAG tag which is used in the exemplification of the invention to determine the association of TRAF
                                                                                                                                          New isolated TRAIL binding protein - which binds to a factor-related apoptosis inducing ligand, used in the treatment of TRAIL-mediated disorders.
                                                                                                                                                                                                   WPI; 1999-095685/08.
                                                                                                                                                                                                                             Walczak H,
                                                                                                                                                                                                                                                                                  26-JUN-1997;
                                                                                                                                                                                                                                                                                                                                       07-JAN-1999.
                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
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                                                                                                                                                                                                                                                                                                            25-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                  WO9900423-A1
                                                                                                                                                                                                                                                                                                                                                                                             Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; TWP-related apoptosis-inducing ligand binding prote
TRAIL-BP; tumour necrosis factor; T cell death; HIV; gene
thrombotic microangiopathy; thrombotic thrombocytopenic pu
haemolytic-uraemic syndrome; systemic lupus erythematosus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human TRAIL-BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAW94672;
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                                                                                                                                                                                                                                                                                 97US-00883529
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein; clotting;
gene therapy;
                                                                                                                                                         tumour necrosis diagnosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                purpura;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
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The present sequence represents a Flag peptide from the present invention. The present invention describes human tumour necrosis factor (TNF) related apportosis-inducing ligand (TRAIL) binding protein (BP). TRAIL-BP can be used for inhibiting the biological activities of TRAIL or for purifying TRAIL. TRAIL-BP proteins can be used for treating a TRAIL-mediated disorder such as T cell death in HIV-infected patients. They can be used for treating thrombotic microangiopathies such as thrombotic

purpura,

syndrome,

of small

Disclosure; Page 3; 47pp; English

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RESULT 52
AAY06471
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                                                                                                                 This peptide comprises an epitope tag that can be used in methods of the invention. The invention discloses a powerful new system for engineering antibody affinity and specificity, by constructing a microbial analogue of the mammalian system's B cell repertoire. Antibodies are displayed on the surface of yeast cells by genetic fusion with yeast cell wall proteins, especially agglutinin proteins. After mutation, variants are selected on the basis of improved binding characteristics with fluorescently labeled targets. The selection method also identifies proteins with enhanced phenotypic characteristics, proteins that are displayed at higher levels, proteins that are secreted at higher levels, proteins that are secreted at higher levels.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           blood vessels or systemic lupus erythematosus. The TRAIL-BP nucleic acids can also be used for gene therapy. They can also be used as carriers for delivering attached agents to cells bearing TRAIL
                                                                                                                                                                                                                                                                                             proteins,
                                                                                                                                                                                                                                                                                                        Selecting
                                                                                                                                                                                                                                                                                                                                   WPI; 1999-430619/36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Epitope tag; anti protein library;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Epitope tag.
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                                                                                             Sequence 8 AA;
                                                                                                                                                                                                                                                                     Disclosure; Page 7; 116pp; English.
                                                                                                                                                                                                                                                                                                                                                          Wittrup KD,
                                                                                                                                                                                                                                                                                                                                                                                                         20-JAN-1998;
26-AUG-1998;
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                                              Local Similarity
nes 8; Conser
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8; Conserv
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                       DYKDDDDK 8
                                                                                                                                                                                                                                                                                            proteins with enhanced phenotypic properties than wild-type is useful for highly specific cancer diagnosis and therapy.
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                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antibody engineering; ary; peptide library.
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98US-00140084.
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                                                         100.0%;
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Pred. No.
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Pred. No.
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                                               Mismatches
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                                                         2.1e+06;
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                                                                      DB 2;
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RESULT 53
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                                                                                                                                                                   RESULT 54
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Matches 8
                                                                                                                                                                                                                                                                                                                                              The present invention describes human tumour necrosis factor (TNF)-related apoptosis-inducing ligand (TRAIL) receptors (TRAILA). TRAILR proteins can be used for binding TRAIL, e.g. to measure or inhibit the biological activity of TRAIL. TRAILR proteins can be used for treating thrombotic microangiopathies, e.g. thrombotic thrombocytopenic purpura (TTP) or haemolytic-uraemic syndrome (HUS), clotting of small blood vessels in e.g. ALDS, multiple sclerosis or systemic lugus erythematosus or for reducing TRAIL-mediated death of T cells in HIV-infected patients. They can also be used to purify TRAIL or TRAIL-expressing cells or as carriers for delivering agents to cells bearing TRAIL. The present sequence represents a Flag peptide form the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New isolated TRAIL receptor polypeptides - used to develop products for treating e.g. thrombotic microangiopathy, multiple sclerosis, systemic lupus erythematosus or HIV infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human, TRAIL; TRAIL receptor; immunoreactive; thrombotic microangiopathy; HIV infection; tumour necrosis factor related apoptosis inducing ligand; TNF related apoptosis inducing ligand; systemic lupus erythematosus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Flag
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           Tumour necrosis factor-gamma; TNF-gamma; human; inflammation; cancer; tumour; metastasis; graft versus host disease; drug screening; therapy;
                                                                                                                                                                                                                                                                                                                           Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 13; 51pp; English.
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 diagnosis; FLAG
                                                       FLAG peptide
                                                                                   26-APR-1999
                                                                                                              AAW88371;
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Pred. No.
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RESULT 55
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                                                                                                                                                                                                                                                                                                                                                                                                                                         This is a FLAG peptide recognised by monoclonal antibody anti-FLAG M2. DNA encoding the FLAG peptide can be ligated to DNA encoding a human serum albumin leader sequence (see AAW88370) and the C-terminal portion (see AAW88369) of novel human tumour necrosis factor gamma (TNFG, see AAW88368) in order to provide a construct for the expression of secretable TNGF proteins in mammalian host cells. Such soluble polypeptides can be used to treat deficiencies of TNFG and disease conditions ameliorated by TNFG. Antibodies, antagonists and inhibitors the polypeptide may be used to treat TNFG-associated diseases, tumours metastases, and to screen for, diagnose and monitor conditions attributable to TNFG, especially inflammation, cancer and graft versus host disease
  Billing-Medel
Granados EN,
                                                                                                                                                             BL210; urinary tract; cancer; tumour; metastasis; diagnosis; therapy; human; epitope; antigen; antibody; FLAG.
                                                                                                                                                                                                 FLAG epitope
                                                                                                                                                                                                                                              AAW97660
                                                                                                                                                                                                                                                                    AAW97660
                                                      08-JUL-1997;
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                                                                                                   21-JAN-1999
                                                                                                                      WO9902734-A1
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                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                  (ABBO ) ABBOTT LAB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 43; Page 70; 106pp; English
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8; Conserv
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 PA, Cohen
Hodges SC,
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                                                      97US-00889866
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                                                                                                                                                                                                                                                                  peptide;
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tion, cancer and graft versus host disease
 M, Colpitts
Klass MR, I
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Pred. No. 2.1e+06;
; Mismatches 0;
Kratochvil JD,
          Friedman PN,
                                                                                                                                                                                                                                                                                                                                                                                      Length 8;
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Russell JC;
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RESULT 56
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This invention describes novel human BS274 protein clones which used as markers for breast cancer and have antitumour activity. also be used as a marker of cancer and its metastases, atypical hyperplasia, fibroadenoma and cysts, so detection/quantification encoded polypeptides and BS274-specific antibodies are used for diagnosis, staging, monitoring, prognosis, in vivo imaging of, a
                                                                                                                                                                                   Billing-Medel PA,
Granados EN, Hodo
                                                                                                                                                                                                                                                                                                                                                                                 Human; breast tissue; BS274; detection; disease; marker; antitumour; breast cancer; metastases; atypical hyperplasia; fibroadenoma; cyst; diagnosis; therapy; antagonist; cytotoxin; immunoassay; drug screen; chromosomal anomaly; immunisation; antisense; triplex; ribozyme.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This peptide comprises a FLAG epitope. An antisense primer incorporating a sequence encoding FLAG has been used in the PCR amplification of human urinary tract specific BL210 cDNA clones (see also AAX07044-57) for the production of a BL210 expression plasmid. The BL210 protein product (see also AAW97655) was recovered from transfected mammalian cells using anti-FLAG monoclonal antibody M2. The invention provides BL210 cDNA clones and polypeptides useful for developing products for detecting, diagnosing, staging, in vivo imaging, preventing or treating diseases or conditions of the urinary tract, including cancer
                                                                                           Example
                                                                                                                     New
                                                                                                                                            WPI; 1999-347484/29.
                                                                                                                                                                                                                                                                             17-NOV-1998;
                                                                                                                                                                                                                                                                                                                                WO9925850-A1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human BS274 protein
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                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
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                                                                                                                  BS274 nucleic acid
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                                                                                        Page 102; 105pp; English.
                                                                                                                                                                     Stroupe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    derived from the BL210 gene - used for diagurinary tract disease, particularly cancer
                                                                                                                                                                                Hodges
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es SC,
pe SD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 fragment
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                                                                                                                  as a marker for breast disease
                                                                                                                                                                                 M, Colpitts
Klass MR, I
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Pred. No. 2.1
0; Mismatches
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Kratochvil JD,
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hes 0;
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Roberts-Rapp
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detection/quantification

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                        This peptide comprises a FLAG epitope. DNA encoding this epitope has been inserted into murine JIP-1 cDNA (see AAV65289) between the first 2 codons of the coding region. Recombinant JIP-1 was expressed in bacterial host cells as GST fusion proteins. Experiments demonstrates that JIP-1 (see also AAW81525) specifically binds JNK (c-Jun NH2-terminal kinase) in vivo. JIP-1 polypeptides and nucleic acids are used in the manufacture of medicaments for treating a pathological condition associated with abnormal expression or activity of JNK, such as a neurodegenerative disease (selected from Parkinson's disease and Alzheimer's disease), a disease (selected from Parkinson's disease and Alzheimer's disease), a
                                                                                                                                                                                                                                                                                                                                                                                                             c-Jun NH2-terminal kinase neurodegenerative disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              JIP-1; JNK-interacting protein; c-Jun NH2-terminal kinase; signal transduction; inhibitor; mouse; neurodegenerative disease; parkinson's disease; hlacheimer's disease; blood clot; stroke; malignancy; cancer; leukaemia; autoimmune disease; inflammation; apoptosis; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FLAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAW81534 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   determining predisposition to these diseases. Antibodies are useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Davis RJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-NOV-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO9849188-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          therapeutic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         immunoassay reagents and
                                                                                                                                                                                                                                                                                                                           Example 2; Page 48; 95pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (UYMA-) UNIV MASSACHUSETTS
   blood clot, stroke,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              rapeutic antagonists (optionally coupled to a cytotoxin), as unoassay reagents and in competitive drug screens. BS274 nucleic acid its fragments, are used as probes and primers (in diagnostic assays)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               peptide epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           œ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           рукрараж в
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DYKDDDDK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dickens M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FLAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               97US-00819177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98WO-US008513.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         peptide; 8
malignancy, leukaemia, an autoimmune disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .0%;
                                                                                                                                                                                                                                                                                                                                                                                                             (JNK)-interacting protein 1 - used, blood clot, leukaemia, autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.1e+06;
thes 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                disease,
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07-JUN-1999

(first entry)

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RESULT 59
AAW99670
ID AAW99
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AC AAW99
XX
DT 07-JU
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Best Local
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                                                                                                                                                       Query Match
Best Local
                                                                                                                                             Matches
                                                                                                                                                                                                                 The invention relates to novel methods for detecting, diagnosing, staging, monitoring, prognosticating, in vivo imaging, preventing, treating or determining a predisoposition to diseases or conditions of the urinary tract, particularly cancer. The methods use products derived from keratin/cytokeratin, cellular apoptosis susceptibility (CAS) or mat-greatific polynucleotides (AAX32486 to AAX32498). These polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Urinary tract; detection; diagnosis in vivo imaging; cancer; keratin; cytokeratin; cellular apoptosis susceptibility; CAS; mat-8; FLAG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAY06792;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY06792 standard; peptide; 8 AA.
                                                                                                                                                                                    Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                         Billing-Medel PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9914372-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FLAG peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-JUN-1999
                               AAW99670;
                                                  AAW99670 standard;
                                                                                                                                                                                                                                                                                                   Example 12a; Page 69; 109pp; English
                                                                                                                                                                                                                                                                                                                      Detection and treatment of urinary tract disease, particularly cancer.
                                                                                                                                                                                                                                                                                                                                           WPI; 1999-229552/19
                                                                                                                                                                                                                                                                                                                                                                Klass
                                                                                                                                                                                                                                                                                                                                                                                                                  15-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                      15-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-MAR-1999
                                                                                                                                                                                                         tract, especially cancer
                                                                                                                                                                                                                                                                                                                                                                                             (ABBO ) ABBOTT LAB
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                                                                                                                         1 рукорорк 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                         Similarity
                                                                                                                                             8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DYKDDDDDK 8
                                                                                                      DYKDDDDDK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ä,
                                                                                                                                                                                                                                                                                                                                                                Kratochvil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; ilarity 100.0%; Conservative
                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   epitope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                  97US-0058925P
                                                                                                                                                                                                                                                                                                                                                                                                                                      98WO-US019362
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ထ
                                                                                                                                                                                                                                                                                                                                                              Cohen M,
vil JD, F
                                                  peptide;
                                                                                                                                                        100
                                                                                                                                                        Russell
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                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                Friedman PN, Goro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 47; DB 2;
Pred. No. 2.1e+06;
); Mismatches 0;
                                                                                                                                                        Score 47;
Pred. No.
                                                    ₿
                                                                                                                                               Mismatches
                                                                                                                                                        2.1e+06;
                                                                                                                                                                  DB 2;
                                                                                                                                                                                                                                                                                                                                                                Gordon
oupe SD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 8;
                                                                                                                                                                  Length 8;
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                                                                                                                                                Gaps
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tag sequence

SEQ

IJ NO:6.

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RESULT 60
AAW95645
ID AAW95
XX
AC AAW95
XX
DT 08-JU
DT 08-JU
DX
XX
ES124
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                                                                                                                                                                                                                                                                                                  δ
                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes a lipocalin homologue, designated zlipol. CC The lipocalin homologue, zlipol, is specifically expressed in testis and commany gland, particularly breast tumour tissue. Based on this tissue cd distribution, zlipol may be used as a diagnostic for breast carcinomas can as a tool for predicting tumour aggressiveness. Agonists can be used cfor transportation of small hydrophobic molecules either in vivo or in cc vitro, and so are useful in specifically promoting the growth and/or cd development of testis-specific cell lineages in culture. Zlipol can be cused to identify inhibitors. Zlipol proteins can also be used to prepare cc antibodies (which can be linked to toxins), and can serve as immunogens. CC transport and/or stabilise small lipophilic molecules, e.g. to protect from gut pH and digestive enzymes. They can also be used to bind small cf fatty acids in blood or tissues to modulate their biological function, ce.g. to transport retinoids or steroids to receptors, in particular as ct therapy for breast cancer, emphysema and diseases of the skin. They may can also play an important role in reproduction. Other uses include anti-
cc inflammatory responses, and antimicrobial activities. Zlipol nucleic acid sequences may be used for gene therapy to increase or inhibit zlipol call to detect genetic abnormalities. The present sequence represents a cc rand to detect genetic abnormalities. The present sequence represents a cc rand to detect genetic abnormalities. The present sequence represents a content of the sequence which is used in an example from the present invention of the content invention.
                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
                                         Human BS124 peptide fragment.
B$124; breast; cancer; detection; diagnosis; prevention; treatment;
                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New lipocalin homologue designated zlipol - whose expression is restricted to testis and mammary gland tissues, particularly breast tumour tissue, used to, e.g. predict tumour aggresiveness.
                                                                                  08-JUN-1999
                                                                                                                                                          AAW95645 standard; protein; 8 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; lipocalin; testis; breast cancer; emphysema;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 5; Page 61; 94pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1999-167367/14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-AUG-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antimicrobial.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                                                    1 DYKDDDDK 8
                                                                                                                                                                                                                                                                                                                                          Similarity
8; Conserv
                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97US-0054867P
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                                                                                                                                                                                                                                                                                                                                                            100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mammary gland; breast tumour; zlipol;
skin disease; reproduction; anti-inflammatory;
                                                                                                                                                                                                                                                                                                                                          <u>,</u>
                                                                                                                                                                                                                                                                                                                                                            Score 47; DB 2;
Pred. No. 2.1e+0
                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                        2.1e+06;
                                                                                                                                                                                                                                                                                                                                                                           Length 8;
                                                                                                                                                                                                                                                                                                                                          Indels
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RESULT 61
AAY31721
XX PRINT PRI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Fibrinogen domain related protein; FDRG; TANGO 115; angiogenesis; adipocyte; diabetes; obesity; inflammation; gene therapy; diagnosis; human; Flag epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The sequence is that of a BS124 peptide fragment. It is detecting, diagnosing, staging, preventing or treating, predisposition to diseases or conditions of the breast,
                     WPI; 1999-540859/45
                                                                      Holtzman DA,
                                                                                                                                                                                                                                                  02-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Flag epitope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Billing-Medel PA, Cohen M, Colpitts TL, Friedman Granados EN, Hodges SC, Klass MR, Kratochvil JD, Scheffel CP, Stroupe SD, Yu H;
                                                                                                                         (MILL-)
                                                                                                                                                                                                  02-MAR-1998;
                                                                                                                                                                                                                                                                                                        10-SEP-1999.
                                                                                                                                                                                                                                                                                                                                                         WO9945135-A1
                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAY31721;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAY31721 standard; peptide; 8 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New isolated BS124 polynucleotides and polypeptides - used for detecting diagnosing, preventing or treating diseases or conditions of the breast, such as breast cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 11a; Page 100; 125pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1999-105623/09.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ABBO ) ABBOTT LAB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-DEC-1998.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 100 hes 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 DYKDDDDK 8
                                                                                                                         MILLENNIUM BIOTHERAPEUTICS INC. DANA FARBER CANCER INST INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DYKODODK
                                                                                                                      FARBER CANCER INST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                      Spiegelman BM,
                                                                                                                                                                                                    98US-00033539
                                                                                                                                                                                                                                                  99WO-US004548
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                                                                      Yoon CH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 47; DB 2;
Pred. No. 2.1e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Friedman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PN, Gordon J; Russell JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            such as breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     determining
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RESULT 62
AAX31944
ID AAX31
AC AAX31
AC AAX31
AC AAX31
AC AAX31
AC AAX31
XX Rad1
XX Rad1
KW Rad1
KW Cell1
CC Frote
PT invo
PT Cycl1
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PT Cycl1
XX This
CC Frote
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence represents the Flag epitope. A human FDRG Flag epitope-tagged protein was constructed by PCR amplification of FDGR cDNA (see AAX87858) using a primer that includes the epitope sequence, ligation into expression vector pMET stop, and expression in HEK 293T cells. FDRG is a secreted molecule that is believed to modulate angiogenesis and to mediate the contribution of adipocytes to various metabolic processes. The invention provides FDRG proteins, antibodies, nucleic acid, and transgenic animals. Also provided are methods for identifying compounds that modulate FDRG activity or which modulate a metabolic disorder involving FDRG. Claimed methods of treating obesity or diabetes, and of regulating angiogenesis, involve administering a small molecule modulator of FDRG, a FDRG nucleic acid molecule or FDRG antibody
This peptide represents the FLAG epitope tag that was used in fusion proteins with human Rad17 (see AAY31937) for expression in mammalian cells, permitting immuno-detection and immuno-purification. Rad17 is novel cell cycle checkpoint control protein. The invention provides 1 polynucleotides and polypeptides, expression constructs, host cells, methods for producing Rad17 polypeptides, antibodies, methods for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rad17; cell cycle checkpoint; human; cell proliferation; immune disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FLAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAY31944 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New isolated Fibrinoge products for treating inflammation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9949043-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                therapy; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-DEC-1999
                                                                                                                                                                                          Novel human RAD17 gene and protein useful for identifying factors involved in cell cycle checkpoint control, e.g. for treatment of
                                                                                                                                    Example 3; Page 28;
                                                                                                                                                                                                                                                                                                                           (ICOS-) ICOS CORP.
(MEDI-) MEDICAL RES COUNCIL
                                                                                                                                                                                                                                                                                                                                                                                       27-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                              29-MAR-1999;
                                                                                                                                                                                                                                                                                           ¥
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             epitope tag
                                                                                                                                                                          control disorders such
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              _
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8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6; Page 93; 152pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DYKDDDDK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DYKDDDDK 8
                                                                                                                                                                                                                                                                                         Herendeen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ĀΑ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fibrinogen Domain Related polypeptides, used to develop treating e.g. obesity, diabetes, cancers, angiogenesis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first
                                                                                                                                                                                                                                                                                                                                                                                       98US-00049760
                                                                                                                                                                                                                                                                                                                                                                                                                              99WO-US006782
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    immune
FLAG ep
                                                                                                                                    75pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  epitope
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    modulator; screening;
cirrhosis; rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.1e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cancer;
arthritis;
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RESULT 63
AAW92756
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Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           identifying binding partners of the polypeptides, and methods of screening for modulators of Radl7 activity. Modulators may be therapeutically useful in treatment of a wide range of diseases, such cancer, in which radl7 activity is involved. They may also be useful chemotherapy and radiotherapy adjuvants
                                                                               The invention provides polynucleotide sequences from urinary tract tissue library BL172. The BL172 polynucleotides (ARX01683-87) are used to detect target BL172 polynucleotides and BL172 mRNA and the BL172 polypeptides are used to detect BL172 antigens/antibodies in a test sample. The identification of certain markers in these methods are indicative of the presence of urinary tract disease, especially cancer. Additionally, the polypeptides are used to detect antibodies that bind specifically to a BL172 epitope derived from a BL172 polypeptide. Host cells containing a recombinant expression vector comprising the BL172 polynucleotide sequences are used to produce the polypeptides containing BL172 epitopes. The methods aid in diagnosis, detection, staging, monitoring.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                         New purified polynucleotide BL172 derivatives and encoded polypeptides - useful in the identification of markers, which are indicative of urinary
                                                                                                                                                                                                                                                                                                                                                                                              Billing-Medel
Granados EN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW92256 standard; peptide; 8 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         chemotherapy and
                                                    The methods aid in diagnosis, detection, staging, monitoring, prognostication, in vivo imaging, prevention and treatment of diseases conditions associated with BL172, especially urinary tract cancer. The
                                                                                                                                                                                                                                                                                                                                                                                 Russell JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BL172; urinary tract; marker; cancer; recombinant; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FLAG peptide epitope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-APR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAW92256;
                                                                                                                                                                                                                                                               Example 11a; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO9855656-A1
                           expression
                                                                                                                                                                                                                                                                                                                                                                                                                                         (ABBO ) ABBOTT LAB
                                                                                                                                                                                                                                                                                            diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       8;
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                                      sequences represents a FLAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DYKDDDDDK 8
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                          plasmid
                                                                                                                                                                                                                                                                                                                                                                               lel PA, Cohen V, Hodges SC, Stroupe SD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                           PA, Cor
Hodges
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                                                                                                                                                                                                                                                               68; 114pp; English.
                                                                                                                                                                                                                                                                                           conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%;
                                                                                                                                                                                                                                                                                                                                                                               M, Colpitts TL, Friedman
Klass MR, Kratochvil JD,
Yu H;
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Pred. No. 2.1
D; Mismatches
                                         peptide
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                                          epitope used
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                                                                                                                                                                                                                                                                                                                                                                                                PN, Gordon J;
Roberts-Rapp
                                            in a BL172
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Best Local Similarity

100 . 0**%** 0**%**;

Score Pred.

47; No.

DB 2; 2.1e+06;

Length

Sequence 8

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ARBSULT 65
AAW94250
ID AAW94
XX AAW94
XX AAW94
XX IS-AE
DT 15-AE
XX XX
DE FLAG
XX PLAG
XX P35 F
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AAY07740
                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                   Query Match
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p35 protein; cowpox; vaccinia; variola; secreted protein; chemokine;
                                                                                                                                                                                                                                                                                                                Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                       This invention describes a novel human breast-specific protein BS200. This protein and its encoding nucleic acids are useful for detecting, diagnosing, staging, monitoring, prognosticating, preventing or treating, or determining predisposition to diseases or conditions of the breast,
                             FLAG peptide
                                                            15-APR-1999
                                                                                         AAW94250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Billing-Medel PA, Co
                                                                                                                   AAW94250 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 11a; Page 119; 124pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             diagnosing, staging, preven breast, e.g. breast cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New breast specific gene BS200 - used to develop products for detecting, diagnosing, staging, preventing or treating diseases or conditions of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-120915/10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-JUL-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human breast-specific BS200 derived protein fragment 1.
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                                                                                                                                                                                                                                                    Similarity
8; Conserv
                                                                                                                                                                                                                                                                                                                                            breast
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             prevention;
                                                                                                                                                                                                                                                       Conservative
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                              epitope
                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                            cancer
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                                                                                                                   peptide; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cohen M, Colpitts TL,
es SC, Klass MR, Krat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein; 8
                                                                                                                                                                                                                                                                100.0%;
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                                                                                                                                                                                                                                                                                   .0%;
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                                                                                                                                                                                                                                                                   Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ₽
                                                                                                                                                                                                                                                     Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   % TL, Friedman PN, Gordon J;
Kratochvil JD, Russell JC;
                                                                                                                                                                                                                                                                                 DB 2;
                                                                                                                                                                                                                                                                  .1e+06;
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RESULT 66
AAW82735
ID AAW8
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention provides nucleic acid sequences (AAX06698-700) encoding p35 proteins of cowpox, vaccinia and variola viruses (AAM94242-44) respectively. p35 is a secreted protein that can be produced by usual recombinant techniques. These p35 proteins can bind to a number of chemotaxis-stimulating cytokines (chemokines). The p35 proteins can be used to treat conditions mediated by chemokines such as monocyte chemotactic protein (MCP)-1, MCP-3, Rantes, Botaxin, macrophage inflammatory protein (MCP)-1 alpha, and MCP-1 beta. They can be used to treat inflammation such as inflammation of the lung, e.g. associated with allergy or asthma. They can also be used to treat inflammation, e.g. psoriasis, or arthritis or autoimmune disorders. The p35 polypeptides can also be used as reagents in assays and for purifying chemokines. The present sequence represents a FLAG peptide epitope used dispractical properties of the invention of the present sequence represents a FLAG peptide epitope used as invention.
               03-DEC-1998
                                                                                                               Chimeric protein; adenoviral fibre protein; trimerisation domain; affinity; substrate; c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Use of poxvirus p35 polypeptides - conditions mediated by chemokines, influenza or autoimmune disorders.
                                                                       Mastadenovirus
                                                                                                                                                                                        29-MAR-1999
                                                                                                                                                                                                                     AAW82735
                                                                                                                                                                                                                                                  AAW82735 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 8 AA;
                                                                                                                                                            Adenovirus chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1999-166574/14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Smith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              chemotaxis; cytokine; monocyte chemotactic protein; MCP-1; MCP-3; MIP-1; Rantes; Eotaxin; macrophage inflammatory protein; inflammation; asthma; lung; allergy; influenza; psoriasis; arthritis; autoimmune disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-DEC-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-SEP-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-FEB-1999
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8; Conserv
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                                                                                                  interaction
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                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                        (first
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95US-00575715
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            of the invention
                                                                                                                                                                                                                                                 peptide;
                                                                                                                                                          protein
                                                                                                                                                                                        entry)
                                                                                                                                                                                                                                                                                                                                                                                                  100
                                                                                                                                                                                                                                                                                                                                                                                                 .0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disorders.
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                                                                                                                                                         FLAG-peptide.
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                                                                                                                                                                                                                                                                                                                                                                                                    Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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e.g. inflammation, asthma, allergy,
                                                                                                                                                                                                                                                                                                                                                                                                    2.1e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                  DB 2;
                                                                                                               gene
                                                                                                                monomer; infection;
gene therapy vector;
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Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This sequence represents a FLAG peptide used in a method to construct novel monomers having an N-terminus of an adenoviral fibre protein and a trimerisation domain. Such monomers have lower affinity for native substrate than the native adenoviral fibre trimer. Cell lines containing such monomers are used (i) to propagate adenovirus for use as gene therapy vectors (for in vitro or in vivo applications, (ii) as reagents for studying adenoviral attachment and infection, and (iii) in receptorligand interaction assays. The new viruses produce minimal ectopic infection (they can not infect native host cells) so are safer as vectors and can be engineered for selective targeting to other cells
                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                    cancer;
                                                                                                                                                                                                                                                                                                        Synthetic FLAG octapeptide
                                                                                                                                                                                                                                                                                                                                  01-DEC-1999
                                                                                                                                                                                                                                                                                                                                                           AAY42254;
                                                                                                                                                                                                                                                                                                                                                                                 AAY42254 standard; peptide; 8 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 14; Page 65; 103pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            useful for, e.g. preparing gene therapy infection for in vitro applications.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New adenoviral fibre trimer with reduced binding to native substrate - useful for, e.g. preparing gene therapy vector with minimal ectopic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1999-059848/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Wickham TJ,
Lizonova A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-MAY-1997;
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                                                                                                                                                                                                                  WO9939734-A1
                                                                                                                                                                                                                                                                                Adenovirus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-JAN-1998;
          gene therapy.
                                                                                                                          06-FEB-1998;
10-SEP-1998;
                                                                                                                                                                 05-FEB-1999;
                                                                                                                                                                                          12-AUG-1999
                      Recombinant adenovirus vectors with modified fiber knob loops, useful
                                                                                                  (UABR-) UAB
                                                 1999-539951/45
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                                                                          DŢ,
                                                                                                                                                                                                                                                                  rus; gene therapy; coxsackievirus adenovirus cystic fibrosis; muscular dystrophy.
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8; Conser
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                                                                                                  RES FOUND
                                                                         Krasnykh VN,
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Yonehiro G;
                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INC.
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98US-0071668P.
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                                                                                                                          98US-0073947P
98US-0099801P
                                                                                                                                                                 99WO-US002549
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
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                                                                          Dmitriev
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 47; DB 2;
Pred. No. 2.1e+06;
; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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                                                                                                                                                                                                                                                                                  receptor; CAR;
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RESULT 68
AAY49544
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This sequence represents a synthetic FLAG octapeptide. DNA encoding this sequence was cloned into the sequence encoding the HI loop of the adenovirus fibre protein knob domain, which was then used in the construction of plasmids encoding a fibre protein. Recombinant adenovirus genomes were generated by homologous DNA recombination in E. coli, before excision of the newly generated genome for virus rescue. The knob domain of the adenovirus fibre protein mediates the initial binding and recognition of the coxsackievirus and adenovirus receptor (CAR) on the cell surface. The HI loop protrudes from the knob domain and connects beta-strands involved in the formation of the cell binding site.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Recombinant adenovirus vectors are used in a number of gene therapy applications; however, the reliance on the CAR means that in certain situations, recombinant virions are sequestered by high CAR-expressing non-target cells while the true target cells, if low in CAR, receive little of the therapeutic gene. Studies of fibre proteins with an HI loop modified by insertion of the FLAG octapeptide (a widely used detection and purification tag) suggested that a heterologous peptide sequence at this location would not affect the structure of the protein and that the HI loop was still accessible for binding. The infectivity of recombinant virions carrying this modified fibre protein was unaffected
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; PRO201; PRO308; PRO309; Nsp1; Nsp2; Nsp3; SH2 domain; EST; expressed sequence tag; tumour; tumourigenesis; diagnosis; cancer identification; proliferation; neoplastic cell growth.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAY49544 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 14; Page 39; 126pp; English
                                                                                                                                                                                                                                                    23-APR-1998;
22-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FLAG epitope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-JAN-2000
                                                                                             New human polypeptides useful to screen antibodies useful to diagnose and treat
                                                                                                                                                                                   Stewart TA,
                                                                                                                                                                                                                                                                                                       23-APR-1999;
                                                                                                                                                                                                                                                                                                                                       28-OCT-1999.
                                                                                                                                                                                                                                                                                                                                                                       WO9954467-A1
                                                           Example 3; Page 54; 152pp; English.
                                                                                                                                                  WPI; 1999-620728/53.
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                                                                                                                                                                                                                     GENENTECH INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                 Lu Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           for insertion into the N-terminus of Nsp1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                        98US-0113296P
                                                                                                                                                                                                                                                                      98US-0082767P
                                                                                                                                                                                                                                                                                                       99WO-US008847
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             peptide; 8 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 47; DB 2; 100.0%; Pred. No. 2.1e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                for antagonists and produce tumors, e.g. cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     <u>,</u>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
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The present invention describes human proteins designated PRO201, PRO308 and PRO309, (also designated Nsp1, Nsp2 and Nsp3 respectively) which are encoded by cDNA clones DNA30676, DNA40575 and DNA61601. The proteins were

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RESULT 69
AAY06609
ID AAY
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Best Local S
Matches 8
                          This peptide represents a FLAG peptide that is recognised by monoclonal antibody M1 (ATCC HB 9259). FLAG has been used in novel fusion proteins of the invention. Such fusion proteins have epitopes of at least 2 of the autoantigens glutamate decarboxylase (GAD65, see AAY06607), islet cell antigen (IAA, see AAY06608) and preproinsulin (PPINS, see AAY06608), the epitopes being connected via linker peptides. The invention also provides the cDNA encoding the fusion protein, a vector and an Escherichia coli cell encompassing the CDNA. The fusion protein is used in an immunoassay
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 combined with radiation treatment or a cytotoxic or chemotherapeutic agent) to inhibit the growth of tumour cells or to treat e.g. leukaemia and immunologic disorders. The antibodies (especially in growth inhibitory amounts) can also be included with a carrier and optionally second antibody or cytotoxic/chemotherapeutic agent in compositions useful as above. They can be used to detect the proteins in cells, by contacting the cell with the antibody and detecting binding, useful to diagnose tumours in mammals (by contacting the antibody with a tissue sample and detecting complex formation). Such diagnosis is especially useful in mammals suspected of having neoplastic cell growth or proliferation. The present sequence represents a FLAG epitope which was inserted into the N-terminus of Nspl CDNA to create pRK.Nsp.FLAG in an anamous formation of the present sequence represents a FLAG epitope which was an example from the N-terminus of Nspl CDNA to create pRK.Nsp.FLAG in an example from the N-terminus of Nspl CDNA to create pRK.Nsp.FLAG in an example of the North PRESENTATION of the North PRESENTATION of the N-terminus of Nspl CDNA to create pRK.Nsp.FLAG in an example of the North PRESENTATION of the N-terminus of Nspl CDNA to create pRK.Nsp.FLAG in an example of the North PRESENTATION of the N-terminus of Nspl CDNA to create pRK.Nsp.FLAG in an example of the North PRESENTATION of the N-terminus of Nspl CDNA to create pRK.Nsp.FLAG in an example of the North PRESENTATION of the N-terminus of Nspl CDNA to create pRK.Nsp.FLAG in an example of the North PRESENTATION of the N-terminus of Nspl CDNA to create pRK.Nsp.FLAG in an example of the North PRESENTATION of the N-terminus of Nspl CDNA to create pRK.Nsp.FLAG in an example of the North PRESENTATION of the N-terminus of Nspl CDNA to create pRK.Nsp.FLAG in an example of the North PRESENTATION of the N-terminus of Nspl CDNA to create pRK.Nsp.FLAG in an example of the North PRESENTATION of the N-terminus of the North PRESENTATION of the North PRESENTATION of the North PRES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New fusion protein, useful for diagnosing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1999-481070/41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hinkkanen
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                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 4; 27pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EP940470-A2
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insulin-dependent diabetes mellitus; IDDM; immunoassay; diagnosis; FLAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FLAG peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-OCT-1999
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simultaneous detection of autoantibodies related
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8; Conserv
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  insulin-dependent diabetes
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AAW92410
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                                                        Query Match
Best Local
                                               Matches
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                                                                                                                This sequence represents a human interleukin-17 receptor (IL-17R) FLAG peptide. IL-17R polypeptides have immunoregulatory activity. They can be used for inhibiting T cell proliferation, or for inhibiting T cell activation. In particular they can be used for preventing or treating organ or graft rejection, autoimmune disease, allergy or asthma. They calso be used for the prevention or treatment of inflammatory disease in which activated T cells play a role or for inhibiting B cell proliferation or immunoglobulin secretion. The IL-17 polypeptides can also be used as immunogens, reagents in in vitro assays, or as binding agents for affinity purification procedures
                                                                                                                                                                                                                                                                          Isolated interleukin-17 receptor DNA - used to develop products for treating e.g. organ or graft rejection, autoimmune disease, allergy, asthma or inflammatory disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IL-17R; human; interleukin-17 receptor; immunoregulator; inhibitor; T cell proliferation; T cell activation; organ; graft; rejection; autoimmune disease; allergy; asthma; treatment; inflammatory disease; B cell proliferation; immunoglobulin secretion; immunogen.
                                                                                            Sequence 8
                                                                                                                                                                                                                                                                                                                                                   Spriggs MK,
                                                                                                                                                                                                                                                                                                                                                                                                   23-MAR-1995;
07-AUG-1995;
                                                                                                                                                                                                                                                   Example 3; Col 29-30; 25pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human IL-17R
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        dependent diabetes mellitus (IDDM)
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8; Conserv
DYKDDDDK
               DYKODDDK
                                                                                             AA;
                                               Conservative
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95US-00538765.
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                                                                                                                                                                                                                                                                                                                                                   WC,
                                                        .0%; Score 47;
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Pred. No. 2.1e+06;
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                                               Indels
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RESULT 71
AAY06911
ID AAY06
AC Grow
KW Grow
KW Catal
KW Inheum
KW Alzhk
KW auto:
XX Syntl
XX IN-SI
XX 
                                                                                           RESULT 72
AAY23636
ID AAY23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CC The invention relates to a growth factor precursor that comprises B-cell courface binding part, T cell surface binding part, antigen cleavable by a CC catalytic antibody (CAb); and a peptide comprising heavy and light chains CC of immunoglobulin. When the antigen is cleaved the B cell surface part CC can interact with its target. The growth factor precursors are used to cells (by inducing mitogenesis, caused by the growth factor released by specific cleavage). The Ag-specific CAb, and to generate CAb from CC such cells (by inducing mitogenesis, caused by the growth factor released by specific cleavage). The Ag-specific CAb can be directed against, e.g. C tumour necrosis factor (treatment of Fheumatoid arthritis or toxic shock CC syndrome); viral docking receptors (treatment of Amman immune virus, CC hepatitis and influenza infections); tumour-specific antigens; amyloid CC plaque (treatment of Alzheimer's disease or myeloma) or IGE (treatment of allergies such as asthma). CAb may also be used for drug detoxification, CC to treat autoimmune or inflammatory diseases and to eliminate CC environmental or industrial pollutants, such as plastics and petroleum. CC cartesponding nucleic acid in a viral or other gene therapy vector. The CC present sequence represents a FLAG peptide epitope
XXX
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Best Local S
Matches 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FLAG peptide epitope
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                              AAY23636;
                                                                                           AAY23636 standard; peptide; 8 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                factor precursor cleaved by antigen-specific catalytic antibody.
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                     Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 8;
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RESULT 73
AAW86029
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                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
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Class II major l
cytotoxic T celi
                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence represents a peptide used to create fusion proteins with the Epstein-Barr virus BZLF2 proteins of the invention. The BZLF2 protein of the invention. The BZLF2 protein is capable of binding to a beta chain of a Class II major histocompactibility complex antigen to inhibit an antigen-specific response. BZLF2 is useful for inhibiting antigen-specific antibody formation, the proliferation of blood mononuclear mononuclear cells, and cytotoxic T cell responses. BZLF2 is also useful for inhibiting undesirable antigen specific responses, e.g. in the treatment or prevention of asthma; for preventing or treating autoimmune disease; and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cytotoxic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Octapeptide used to make BZLF2 fusion proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-SEP-1999
                                                                                                 UTI16; urinary tract; epitope; antigen; detection; diagnosing;
monitoring; in vivo imaging; cancer; agonist; antibody; tumour;
                                                                                                                                                                                                                                                                                                                                                                                      Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Col 31-32; 25pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1999-418295/35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Spriggs MK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Comeau MR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-APR-1994;
28-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-SEP-1997;
                                                             Synthetic
                                                                                       metastasis.
                                                                                                monitoring;
                                                                                                                                      FLAG peptide
                                                                                                                                                                   23-FEB-1999
                                                                                                                                                                                                                  AAW86029 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Epstein-Barr virus BZLF2 proteins.
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                                     WO9851824-A1
                                                                                                                                                                                                                                                                                                                                                                                                             for preventing tissue or organ transplant rejection
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                                                                                                                                                                                                                                                                                                                                                  Similarity
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Cohen JI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           in; beta chain; blood mononuclear mononuclear cell; jor histocompatibility complex antigen; proliferation; cell response; antigen specific response; asthma; lisease; transplant rejection.
                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                        epitope
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95US-00430633.
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                                                                                                                                                                                                                                                                                                                                      Score 47; DB 2;
Pred. No. 2.1e+06;
); Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                            Length
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RESULT 74
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Best Local S
Matches 8
                                                     WPI; 1999-411830/35.
N-PSDB; AAX89170.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            containing the UT116 gene can be used to produce a UT116 polypeptide recombinantly. This polypeptide has at least one UT116 epitope which can be used in a method for detecting UT116 antigen in a test sample. The polynucleotides and polypeptides are useful for detecting, diagnosing, monitoring, staging, prognosticating, in vivo imaging, preventing, treating or determining the predisposition of a subject to diseases and conditions of the urinary tract, such as urinary tract cancer. Antibodic specifically binding to an epicope of UT116 antigen, and agonists are useful for treating urinary tract diseases, tumours and metastases. The present sequence represents a FLAG peptide epitope used in the construction of plasmids for the expression of UT116 polypeptide
  New vertebrate slit protein -
                                                                                                                                                                                                                                 05-DEC-1997;
                                                                                                                                                                                                                                                                                                                                 JP11164690-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                         muscle; endocrine
                                                                                                                                                                                                                                                                                 22-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Vertebrate-derived protein; slit protein; diagnosis; cancer; nerve;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Peptide encoded by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAY27153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAY27153 standard; peptide; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a method of detecting the presence of a target UT116 polynucleotide in a test sample using UT116 gene-specific sequenc (AAV80386 to AAV80397). Host cells transfected with an expression vecto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New method for detecting diseases of the urinary tract - a UTI16 polynucleotide, protein or antibodies, used for protein or antibodies.
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Hodges SC, Klass
                                                                                                                               (ASAH ) ASAHI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 11; Page 67; 113pp; English.
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8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polynucleotide, protein or ant
g urinary tract infections and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DYKDDDDDK 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                      system.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seq ID No:
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R, Kratochvil JD,
                                                                                                                                  KOGYO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100
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useful for diagnosis and treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 of JP11164690
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Roberts-Rapp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cancer
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ches 0;
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    comprises
    preventing

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Matches
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    This peptide comprises a FLAG epitope. An antisense primer incorporating a sequence encoding FLAG has been used in the PCR amplification of human breast specific BS202 cDNA clones (see also AAX07004-09) for the production of a BS202 expression plasmid. The BS202 protein product (see also AAW97643) was recovered from transfected mammalian cells using anti-FLAG monoclonal antibody M2. The invention provides BS202 cDNA clones and polypeptides useful for developing products for detecting, diagnosing, staging, preventing or treating diseases or conditions of the breast, including breast cancer
                                                                                                                                                                                 diagnosing, staging, preast such as breast
                                                                                                                                                                                                                                                                                        Billing-Medel Granados EN,
                                                                                                                                                                                                           New breast specific gene,
                                                                                                                                                                                                                                                                            Stroupe
                                                                                                                                                                                                                                                                                                                                                                07-JUL-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BS202 protein; breantigen; epitope;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a vertebrate-derived protein containing an amino acid sequence shown in AAY27137 and AAY27139. The vertebrate-derived protein has at least 55 % homology to one of sequences shown in AAY27141-Y27143, and has slit protein-like activity. The vertebrate slit proteins encoding nucleic acid sequences have at least 60% homology to nucleic acid sequences AAX89161-163. The vertebrate-derived proteins can be produced recombinantly by transforming host cells with expression vectors comprising the encoding nucleic acids. The proteins of the invention are for diagnosing and treating cancer of the nerves, muscle and/or endocrine
                                                                                                                                                                                                                                              WPI; 1999-120783/10
                                                                                                                                                                                                                                                                                                                                                                                                                        21-JAN-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FLAG peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW97648;
                                                                                                                                                                                                                                                                                                                                 (ABBO ) ABBOTT LAB.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page 88;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                      Page 70; 110pp; English
                                                                                                                                                                                                                                                                                      PA, Co
Hodges
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            breast cancer; tumour; metastasis; diagnosis; therapy;
                                                                                                                                                                                                                                                                                                                                                              97US-00888894
                                                                                                                                                                                                                                                                                                                                                                                             98WO-US014046
                                                                                                                                                                                                                                                                                      Cohen M, Colpitts TL, Friedman s SC, Klass MR, Kratochvil JD,
                                                                                                                                                                                                 preventing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  102pp; Japanese
                                                                                                                                                                                    cancer
                                                                                                                                                                                                 BS202 - used to develop products for detecting, enting or treating diseases or conditions of the
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Pred. No.
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                                                                                                                                                                                                                                                                                                    Friedman PN,
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                                                                                                                                                                                                                                                                                        Russell JC;
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AAM96302
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XX TANGG
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The murine TANGO-69 cDNA encodes a type II membrane protein, which is the CC murine homolog of the human herpesvirus entry mediator (HVEM) ligand, CC LIGHT. LIGHT is a novel cytckine and is termed LIGHT because it shows CC homolgy to Lymphotoxins, exhibits Inducible expression and competes with CHSV Glycoprotein D for HVEM, a receptor expressed by T-lymphocytes. HVEM CC and TANGO-69 are members of the tumour necrosis factor receptor (TNFR) CC ligand superfamily. The TANGO-69 protein regulates a variety of cellular processes including cell proliferation, programmed cell death and immune responses by binding to a membrane-bound form of HVEM (mHVEM) to inhibit CC mHVEM activity. The protein plays an essential role in inhibiting CC are useful for detecting the polymucleotide in samples (especially CC mRNA) and antibodies directed against the TANGO-69 coding sequence CC are useful for detecting the polymucleotide in samples (especially CC polypeptide are useful for modulating the activity of the polypeptide are useful for modulating the activity of the polypeptide and compuned which bind to the TANGO-69 coding sequence CC polypeptides are useful for interacting with proteins in the TANGO-69 signalling pathway, binding the TANGO-69 receptor, interacting with mHVEM canduction on endothelial cells and coagulation. Two tyckine production, CC leukocyte activation, wiral proliferation, cell death adhesion molecule conduction on endothelial cells and coagulation. Two primers (AAXO8401, AAXO8402) were used to amplify the TANGO-69 sequence for its subcloning cinto the expression vector ppicZalpha A for expression in Pichia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Newly isolated murine useful for regulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1999-205132/17.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C-terminal tagging
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (MILL-) MILLENNIUM BIOTHERAPEUTICS INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        polynucleotide encoding a TANGO-69 polypeptide cell proliferation and differentiation, and cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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Pred. No. 2.1e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
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RESULT 77
AAW95562
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          CC polynuclectide that comprises treating a test sample with at least one CC LS170-specific nucleic acid that has at least 50 percent identity with CC any of the sequences (AAXONSO1-809) which are fragments derived from CC LS170 gene. These LS170 nucleic acid fragments represent a set of CLS170 gene. These LS170 nucleic acid fragments represent a set of CC prevention and treatment of lung disease, specifically cancer, and to CC prevention and treatment of lung disease, specifically cancer, and to CC indicate predisposition to such disease. Particularly detection of LS170 CC indicate predisposition to such disease. Particularly detection of LS170 CC polynucleotide, LS170 antigens, or anti-LS170 antibodies is indicative of CC indicate predisposition of such disease. Particularly detection of LS170 cc polynucleotide containing at least one epitope in their sequences, CC indicate predisposition to such disease procession system comprising the LS170 cc polypeptides containing at least one epitope in their sequences, CC indicated to isolate related sequences; as standards and reagents in CC assays; as targets for drug screening. LS170-related nucleic acid CC can be used to isolate related sequences; as standards and reagents in cassays; as targets for drug screening, and as components or targets for CC used to deliver therapeutic agents to LS170-expressing cells; directly as therapeutic agents (by neutralising LS170-expressing cells; directly as therapeutic agents (by neutralising LS170 polypeptides); in competitive containing drug screens, and to generate anti-iddotypic antibodies for use containing designs. Ab can be contained and drug design. The present sequence represents a FLAG peptide circulary design.
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FLAG peptide epitope used in a LS170 expression plasmid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New LS170 nucleic acid from lung tissue - useful for detecting monitoring, preventing and treating lung disease, particularly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Billing-Medel PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-JUN-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1999-060335/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to the detection of a target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 11; Page 70; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ABBO ) ABBOTT LAB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gene;
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8; Conserv
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ilarity 100.0%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cohen M, Colpitts TL, es SC, Klass MR, Kratoc pe SD;
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Pred. No. 2.1e+06;
); Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cancer; drug screening; FLAG
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RESULT 78
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                                                                                                                                      A method has been developed for screening for substances which inhibit
C the binding of TAKI polypeptide to TABI polypeptide. The method
C comprises: (a) contacting the polypeptide in the presence of a sample;
and (b) detecting the amount of bound polypeptide, in which the sample
C can be pre-mixed with TAKI or TABI polypeptide first. The transforming
G growth factor (TGF)-beta inhibitory substances can be used in drugs for
indications e.g. as TGF-beta signal transmission inhibitors or
activators, or extracellular matrix protein production enhancement
c activators, or monocyte migration inhibitors or activators, or
physiological activity induction inhibitors or activators, or
immunosuppression inhibitors or activators, or amyloid beta protein
c precipitation inhibitors or activators, and such substances can also be
inhibitors of the TAKI polypeptide function, particularly kinase
activity. The present sequence represents a peptide from an example of
the present invention
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Best Local S
Matches
                                               Query Match
Best Local Similarity
Matches 8; Conserv
                                                                                                      Sequence 8
                                                                                                                                                                                                                                                                                                                                                                                      Screening for TGF- beta inhibitory substances, which are useful as drugs for treatment of diseases relating to its disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; TAB1; TAK1; screening; inhibition; TGF-beta; transforming growth factor beta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 8
                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 58; 195pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CHUS ) CHUGAI SEIYAKU KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22-OCT-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FLAG peptide
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                                                   Conservative
                                                                                                                                 invention
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                                                               100.0%;
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                                                               Score 47;
Pred. No.
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Pred. No. 2.1e+06;
Mismatches 0;
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                                                  Mismatches
                                                             DB 2; 1
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DYKDDDDX 8

Conservative

Indels

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Gaps

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CC This sequence represents a FLAG peptide used in the isolation of a a CC novel human beta chemokine, ZCHEMO-8. Altered levels of ZCHEMO-8 may be CC indicative of pathological conditions, including infections, cancer, CC myelopoietic disorders, autoimmune disorders and immunodeficiencies. The CZCHEMO-8 polypeptides can be used, e.g. to reduce the damage in ischaemic CC and reperfusion injuries, in a wound healing regime to stimulate an CC infiltration of immune cells (e.g. monocytes, neutrophils, T lymphocytes or basophils) to a wound site to facilitate healing. ZCHEMO-8 may be used CC for transplants. ZCHEMO-8 polypeptides could be used to further define CC the role of chemokines in mediating suppression of HIV replication in CC CD4+T-cells and limiting progression of HIV replication in CC CD4+T-cells and limiting progression of HIV replication in CC CD4+T-cells and limiting progression of HIV replication in CC CD4+T-cells and limiting progression of HIV replication of therapy, to protect haematopoietic cells. ZCHEMO-8 antagonists may have a beneficial therapputic effect in diseases where the inhibition of CC activation of certain macrophages, neutrophils, basophils, B lymphocytes and/or T cells may be effective. Such diseases include autoimmune CC diseases e.g. multiple sclerosis, insulin-dependent diabetes and systemic CC upus erythematosus, rheumatoid arthritis, allergies, asthma or CC antegronists for chronic inflammatory and infective diseases. Antagonists may be used to dampen or inactivate ZCHEMO-8 during activated immune CC creepings. The products can also be used for detection, diagnosis, drug
Query Match
Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New isolated human beta-chemokine, ZCHEMO-8 - for treating e.g. ischaemia, reperfusion, wour diseases, inflammation, asthma or infections.
                                                                               Sequence 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 10; Page 94; 131pp;
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Similarity
8; Conserv
                                                                                                                 The products or gene thera
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                                                                                                                   ducts can also therapy
100.0%; Score 47; DI
100.0%; Pred. No. 2.1
tive 0; Mismatches
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                   2.1e+06;
                                          DB 2;
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RESULT 80
AAY06900
RESULT 81
AAY25328
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Matches 8
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29-SEP-1997;
19-MAY-1998;
19-MAY-1998;
                                                                                                                                                                                                                                                                                                                              The invention relates to a human secretory protein, ZSIG-11. Host cells containing a vector comprising the ZSIG-11 nucleic acid are used for the recombinant expression of the protein ZSIG-11 is a novel ligand polypeptide and specific antibodies can be used to detect its presence in a biological sample. Probes derived from ZSIG-11 nucleotide sequences can also be used in detection of ZSIG-11 RNA. ZSIG-11 is expressed at high levels in testis, and could be used to identify study prohormone convertasses or endoproteases that exhibit testis specificity. Antagonists, including antibodies, are useful for inhibiting or eliminating the function of ZSIG-11. It is possible that ZSIG-11 and its antagonists will be useful as fertility inducing therapeutics
                                                                                                                                                                                                                                                                                                        Sequence 8 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Secretory protein; ZSIG-11; ligand polypeptide; testis; endoprotease; prohormone convertase; fertility; therapeutic; human; FLAG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FLAG affinity peptide sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Polynucleotide encoding a human secretory protein, ZSIG-II
Pancreatic disease; PA153; human; cytostatic; detection; antigen; anti-PA153; antagonist; therapy; treatment; tumour; metastasis;
                                                                     03-SEP-1999
                                        Human pancreatic PA153 derived peptide.
                                                                                                                          AAY25328
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8; Conserv
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                                                                                                                       standard; protein; 8
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97US-00939897.
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Pred. No.
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RESULT 82
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Granados EN, Hodges
Russell JC, Stroupe
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                      Billiing-Medel PA, Cohen
Granados E, Hodges SC, F
Russell JC, Stroupe SD;
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                                                                                                                      LAB.
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Jes SC, K
Jee SD;
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                                             Klass MR,

 Colpitts TL,
Klass MR, Krat

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                                               Colpitts TL, Friedm
s MR, Kratochvil JD,
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Kratochvil JD, Roberts-Rapp
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                                                  Roberts-Rapp
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RESULT 83
AAY04141
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Best Local
                                                                                                        The present sequence represents a peptide from a human slit-like protein. Slit-like proteins can be used for the prevention and the treatment of diseases in spinal cord, thyroid gland, ovary, prostate, renal gland, small intestine, heart, trachea, thymus, lymph node, muscular system and
                                                                                                                                                                                                         New slit-like polypeptide - useful for prevention and treatment of diseases in spinal cord, thyroid gland, ovary, prostate, renal gland, small intestine, heart, trachea, thymus, lymph node, muscular system and
                                                                         Sequence 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention provides BS106 polynucleotides (AAX55578-82) and polypeptides (AAX13466-470). The BS106 nucleic acids correspond to a breast tissue gene and can be used as markers for breast tissue disease. The products can be used for detecting, diagnosing, staging, monitoring, prognosticating, preventing or treating, or determining diseases or conditions associated with BS106, especially breast cancer
                                                                                                                                                                                                                                                                                                                                       02-SEP-1997;
                                                                                                                                                                        Example
                                                                                                                                                                                                                                                                                       (ASAH ) ASAHI KASEI KOGYO KK
                                                                                                                                                                                                                                                                                                                02-SEP-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human slit 3 peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                ; slit-like protein; slit 3; slit 1; prevention; treatment; disease; l cord; thyroid gland; ovary; prostate; renal gland; intestine; heart; trachea; thymus; lymph node; muscular system;
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                                     Similarity
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ilarity 100.0%;
Conservative (
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                    Score 47; DB 2; 1
Pred. No. 2.1e+06;
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                                                                                                                                          The present invention describes peptides which have a moderate or high CC affinity for each other, when added as extensions to both the N-terminus CC and C-terminus of a protein, can be used to help fold the protein into a CC compact structure. This compact structure is more stable to proteases. A CC composition (A) comprises at least a first dimerisation peptide (I) CC comprising the sequence (S) that is no more than 8 amino acids long, CC where the composition optionally comprises a second dimerisation peptide (I): NH2-X1-X2-X3-X4-X5-COOH (S1) where X1 to X4 = Ala, Val, Ile, Leu, CC [II]: NH2-X1-X2-X3-X4-X5-COOH (S1) where X1 to X4 = Ala, Val, Ile, Leu, CC [II]: The compositions of the invention are displayed intracellularly or extracellularly and are useful to identity binding CC pathways. A library of constrained proteins may be evaluated in vivo for its bioactive potential. The invention can be used to access molecules or CC constrained protein. The compositions of the isolation of the CC constrained protein. The compositions of the invention are useful as a cc scaffold for gene therapy and for potential use as a therapeutic in physiological fluids. The present sequence is used in the exemplification of the physiological fluids. The present sequence is used in the exemplification
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                                                       Query Match
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Matches 8
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library
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      proteins
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   dimerization peptides which self-associate are used with other ins to effect the formation of compact structures.
                    1 DYKDDDDK 8
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                                                         Similarity
8; Conser
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Pred. No.
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RESULT 85 AAE16563

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RESULT 86
AAW97673
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DT 87
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Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    displayability on a yeast cell surface. The method comprises transforming yeast cells with a vector that expresses a test protein fused to a yeast cell wall protein (AGA2), contacting the cells with a label that binds to proteins displayed on the cell wall, and isolating label-bound cells, where the test protein is from a variegated population generated by mutagenesis. The invention is also directed to new processes for engineering T cell receptor for improved binding properties. Improved T cell receptor for improved binding properties. Emproved T autoimmune diseases such as arthritis, diabetes or multiple sclerosis, autoimmune diseases such as arthritis, diabetes or multiple sclerosis. The methods are useful to select proteins with altered affinity altered specificity or conditional binding. The present sequence is an epitope tag fused between protein of interest and yeast cell wall protein AGA2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 40; Col 60; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              proteins,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Selecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Wittrup KD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-MAY-1996;
30-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US6331391-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Yeast cell wall cancer; sepsis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAE16563 standard; peptide; 8 AA.
                                                 MKK3; human; mitogen activated protein kinase kinase; MAP kinase kinase; signal transduction; inflammation; psoriasis; AIDS; cancer; apoptosis; therapy; FLAG; epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (UNII ) UNIV ILLINOIS FOUND
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                                                                                                                                                                  FLAG epitope
  Synthetic
                                                                                                                                                                                                                             10-MAY-1999
                                                                                                                                                                                                                                                                                    AAW97673;
                                                                                                                                                                                                                                                                                                                                        AAW97673 standard; peptide; 8
                                                                                                                                                                                                                                                                                                                                                                                          98
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tag
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DYKDDDDK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DYKDDDDDK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           proteins with enhanced phenotypic properties than wild-type is useful for highly specific cancer diagnosis and therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               #7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  invention relates to a method for selecting proteins for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kranz DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                             (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            fused to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  96US-0018741P
97US-00866398
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autoimmune disease; arthritis; diabetes.
                                                                                                                                                                                                                          entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            yeast cell wall protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kieke M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .1e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 8;
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CC -69) encoding human mitogen activated protein (MAP) kinase kinase 3 (see CC AAW97668) and MKK4 (see AAW97670-72) by insertional overlapping PCR. CC Recombinant MKK polypeptides were used in in vitro phosphorylation CC experiments. MKK3 phosphorylated p38 MAP kinase but not JNK1 or ERK2. CC the transduction of specific signals from the cell surface to the nucleus CC along specific pathways. They are useful for screening reagents which CC modulate MKK activity. Such agents can be used to prevent or treat MKK-CC mediated disorders, e.g. inflammation, oxidative damage or stress-related CC proliferative disorders, e.g. psoriasis, AIDS, malignancies of e.g. the CC skin, bone marrow, lung, liver, breast, gastrointestinal system and CC genito-urinary tract. Agents which inhibit the activity or expression of CC MKK inhibit cell growth or cause apoptosis. MKK7 polynucleotides (see CC AAX07059-64) and polypeptides (see AAW97662-67) are claimed
                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New isolated mitogen-activated protein kinase kinase isoforms - used develop products for treating e.g. inflammatory disorders, oxidative damage, proliferative disorders or autoimmune disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-JUL-1998;
          07-MAY-1996;
05-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This peptide comprises a FLAG epitope. DNA coding for the epinserted between codons 1 and 2 of cNA clones (see AAX07065
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 3; Page 31; 168pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-120771/10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9902547-A1
                                                                                                                                                                                                                       FLAG peptide epitope tag sequence.
                                                                                                                                                                                                                                                      03-MAY-2000
                                                                                                                                                                                                                                                                                   AAY77538;
                                                                                                                                                                                                                                                                                                               AAY77538 standard; peptide; 8 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 inserted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UYMA-) UNIV MASSACHUSETTS
                                                        16-DEC-1997;
                                                                                     11-JAN-2000
                                                                                                                                             Synthetic.
                                                                                                                                                                                         Neurotactin;
                                                                                                                 US6013257-A
                                                                                                                                                                         neuroprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                   1 рукрараж в
                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                         рукрарак в
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                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
nilarity 100.0%;
Conservative (
                                                                                                                                                                           multiple sclerosis; chemokine; inflammation; brain; human; ive; antiinflammatory, cytostatic; antibacterial; FLAG.
                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              97US-00888429
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          96US-00643798
97US-00851160
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 47; DB 2;
Pred. No. 2.1e+06;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ç
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA coding for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     epitope was
                                                                                                                                                                                                                                                                                                                                                                                                                                                     0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             - used to
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RESULT 88
AAB03362
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Best Local
         Various mediators are known to cause pain and inflammation e.g. prostaglandins, histamine, bradykinin and serotonin. The present invention relates to inhibition of pain and inflammation at a wound during surgery by delivering locally and perioperatively to the site a solution containing at least one soluble receptor antagonist. The solution would inhibit locally the mediators of pain, inflammation, spand restensis. The present sequence is the N-terminal sequence of a
                                                                                                                                                      Preemptive inhibition of pain and inflammation in surgical procedures comprising application of soluble receptors, e.g. TNF, optionally with pain and inflammation inhibitor, directly at the wound site, useful e.g.
                                                                                                                     Disclosure; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                          Surgical irrigation solution; anti-inflammatory; anti-pain; anti-spasm;
                                                                                                                                                                                                                                           Demopulos
                                                                                                                                                                                                                                                                                                05-NOV-1998;
                                                                                                                                                                                                                                                                                                                        05-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                  11-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention provides a method of treatment of multiple sclerosis that comprises administering an antibody that binds to human chemokine neurotactin. The antibodies inhibit neurotactin, a chemokine that is associated with inflammation, particularly in the brain. The antibodies associated with inflammation particularly in the brain. The antibodies are useful for treatment of multiple sclerosis. The present sequence
                                                                                                                                                                                                                                                                     (OMER-)
                                                                                                                                                                                                                                                                                                                                                                                                                                  anti-restenosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAB03362
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        represents a FLAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               proinflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-146535/13
                                                                                                                                                                                                                 2000-365358/31.
                                                                                                                                                                                                                                                                    OMEROS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 рукророк в
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       multiple sclerosis by administering
                                                                                                                                                                                                                                           GA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        рукорорк в
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequence of a soluble human IL-1R fusion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                           Palmer
                                                                                                                                                                                                                                                                    MEDICAL
                                                                                                                                                                                                                                                                                               98US-0107256P
                                                                                                                                                                                                                                                                                                                        99WO-US026330
                                                                                                                                            dentistry.
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                                                                                                                  70; 122pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                analgesic; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       peptide epitope tag
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    peptide; 8
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                                                                                                                                                                                                                                          PР,
                                                                                                                                                                                                                                                                     SYSTEMS
                                                                                                                                                                                                                                           Herz JM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             neurotactin.
                                                                                                                     English.
 receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 47; DB 3;
Pred. No. 2.1e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                               IL-1R fusion protein; interleukin-1.
fusion protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          an antibody that
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein
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Best Loc Matches Query Match

Local

Similarity 8; Conser

100.0%; ilarity 100.0%; Conservative 0;

0

Score 47; DE Pred. No. 2.1 0; Mismatches

2.1e+06;

DB 3;

Length 8;

Indels

0

Gaps

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RESULT 89
AAY99877
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                   The present sequence is the FLAG epitope which may be fused to antibodies as a label for their identification and isolation. The label may be added to monoclonal antibodies which bind to conformational epitopes of the envelope glycoprotein E2 from multiple HCV genotypes. These antibodies may be useful in the prevention and treatment of the majority of HCV infections. For example, a subset of antibodies (CBH-2, CBH-5, CBH-7, CBH e0. CBH-8B and CBH-11) have the ability to prevent the binding of HCV E2 proteins of multiple genotypes to human CDB1, a possible coreceptor for HCV infection. Thus the antibodies are useful in passive immunotherapy strategies for reducing viral load of infected individuals and interfering with infection of target cells
                                                                                                                                                                                                                                                Detection of hepatitis C virus (HCV) infection comprises using monoclonal antibodies binding to conformational epitope of an envelope protein from more than one HCV genotype, the monoclonal antibodies are also used as a
                                                                                                                                                                                                                                                                                                                                      Foung
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                proinflammatory and it is thought that IL-1 effects would be prevented blocking IL-1 binding to IL-1R, e.g. IL-1 induced inflammation and cartilage destruction within joints. The IL-1R fusion protein consists the IL-1R binding domain and binds to IL-1 with high affinity. Local delivery of an IL-1R soluble receptor, such as the IL-1R fusion protein in an irrigation solution during joint surgery may be useful as a
                                                                                                                                                                                                     Disclosure; Page 16;
                                                                                                                                                                                                                                                                                                                                                                                                 05-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                              29-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200026418-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FLAG epitope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY99877 standard; protein; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cartilage protective agent
                                                                                                                                                                                                                                                                                                                                                                    (STRD ) UNIV LELAND STANFORD JUNIOR.
                                                                                                                                                                                                                                                                                                                                      SKH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hepatitis C virus; HCV; envelope glycoprotein e; immunotherapy; antibody; CBH.
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                                                                                                                                                                                                                                                                                                                                        Hadlock KG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%;
ilarity 100.0%;
Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antibody label.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                               98US-00187057
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                                                                                                                                                                                                                                    CV genotype,
HCV.
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                                                                                                                                                                                                     85pp; English.
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hes 0;
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                                                                                                                                                                                                                                               pe protein
also used
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                               protein, where the sequence region contains the alpha-secretase cleavage site at a reference of 6 residues at the N-terminal and 12 residues at the C-terminal; and (b) a gene for a protease protein (PP), that either comprises a proteolytically active necessary sequence region or a sequence region of the disintegrin metalloprotease ADAM 10 from a cow (Bos taurus), from a human or other mammal or a mutant of this, which shows the same enzymatic properties, where the genes are under the control of heterologous promoters; or a heterologous RNA coding for a SP and a PP. The products of the invention have nootropic and neuroprotective activity and can be used for gene therapy. The protease CC substrate proteins, especially human amyloid precursor protein. Dominant comparise forms of bovine, human or other mammalian disintegrin. CC metalloprotease ADAM 10 proteins and their coding sequences are useful for suppressing the alpha-secretase activity of a cell. Nucleic acid constructing the proteins and recombinant cells are useful for identifying secretases and pharmaceutical agents and to identify risk constructing secretases and pharmaceutical agents and to identify risk constructions.
                                                                                                                                                                                                                                                                                                                                                                                                                    nucleic acids encoding a region of human amyloid precursor protein containing an alpha-secretase cleavage site and a protease or a heterologous RNA coding for a substrate protein and a protease. The invention also describes a recombinant cell, characterized in that it contains recombinant nucleic acids comprising either: (a) a gene for a substrate protein (SP), which comprises a sequence region of 18 amino acids of the human amyloid precursor protein (APP) or a homologous protein where the general recombination of the human amyloid precursor protein (APP) or a homologous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        amyloid precursor protein and alpha-secretase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Recombinant cells, for identifying alpha-secretase active agents and identifying risk factors associated with Alzheimer's disease, comprise
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-588391/56.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fahrenholz F,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   disintegrin-metalloprotease; protease;
gene therapy; Alzheimer's disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes a novel recombinant cell comprising recombinant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 4; Page 8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APP; amyloid precursor protein; human;
Sequence 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    epitope tag
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              alpha-secretase; ADAM 10;
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RESULT 91
AAY87937
ID AAY8
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                                                         Matches
                                                                                    Query Match
                                                                                                                                                                                        This invention describes a novel peptide (P), homologous to the intracellular C-terminal domain of a mammalian HDL-receptor. The products of the invention can modulation cholesterol, bile acid and lipid metabolism. The peptides may alter the activity of the enzymes in the peroxisome to inhibit cholesterol synthesis and promote cholesterol elimination. The cDNA encoding the peptide is used in gene therapy and is also used for the production of the peptide. The peptide can be used in the treatment of hyperlipidemia and for the modulation of cholesterol, bile acid and/or lipid metabolism of a patitent. The peptide can also be used for the identification of drug targets for treatment of hyperlipidemia. The drug targets thus identified are targets for modulation of the metabolism of cholesterol, bile acid and/or lipids. In modulation of the metabolism of cholesterol, bile acid and/or lipids. In
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Carlsson B, Carlsson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     treatment; hyperlipidemia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Intracellular C-terminal HDL-receptor domain FLAG epitope #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-SEP-2000 (first entry)
                                                                                                                                             addition the peptide may also be used for the production of a medicinal product for treatment of hyperlipidemia. This sequence represents a FLA epitope used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Page 8; 23pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                     treatment of hyperlipidemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-OCT-1998;
                                                                                                                   Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                     mammalian HDL-receptor useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                 New peptide homologous to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (JOHN/) JOHNSON M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CARL/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  intracellular C-terminal HDL receptor domain; cholesterol; bile acid;
                                                                         Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   density lipoprotein; lipid metabolism; peroxisome; gene therapy;
                            1 рукрарак в
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                                                                         Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                     intracellular C-terminal domain of a in identifying drug targets and the
                                                                         Score 47;
Pred. No.
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                                                                          2.1e+06;
                                                                                          DB 3;
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                                                                                        Length 8;
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Best Local Similarity

100.0%;

Score 47; Pred. No.

DB 3; 2.1e+06;

Length 8;

Query Match

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DYKDDDDDK 8

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AAB10094
ID AAB1
XX AAB1
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XX Synt
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PR 30-\
PF 14-I
PR 30-\
EN WPI
XX Dis
PX WPI
PA (DB)
PA (BD)
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                                                                                                                                                                                                                                        CC especially monomers for combinatorial synthesis of molecule libraries, to CC a carrier by embedding (I) in a matrix, an applying the matrix to the CC carrier either as a solid or a gel. (I) is first embedded in a matrix (CC comprising at least 1 first solvent (SI) that is solid at below 90 CC (preferably 50) plusoC so as to form transport units (TU) that can be CC moved individually. TU are applied to the carrier as solid, at below 90 CC plusoC or are applied dissolved in second solvent (S2), which at the same CC solid or gel is formed on the carrier. (I) is then rendered mobile by CC altering the physical conditions, brought into contact with the carrier CC the surface by a physical process and coupled covalently to molecules (II) on CC the surface (or it undergoes, or catalyzes, a reaction with (II). Coupled (I) may be (or may generate) many different substances and application of CC (I) can be repeated sequentially, with intermediate rinsing off of any CC uncoupled (I). The method is used to produce combinatorial chemical CC identifying lead compounds (specific binding agents) for drug diseases of unknown or uncertain etiology) in blood samples or for CC diseases of unknown or uncertain etiology) in blood samples or for CC development. Encapsulation of (I) significantly reduces its diffusion CC compact assembly of a library relatively free of artefacts. The library CC may have a very high density, increasing the chances of detecting CC simultaneous screening for many different diseases. This sequence CC simultaneous screening for many different diseases. This sequence CC simultaneous screening for many different diseases. This sequence
                                                              Query Match
Best Local S
Matches 8
                                                                                                                                                                                                  Sequence 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Applying materials to a support, useful for combinatorial synthesis of peptides, e.g. for diagnosis, by incorporating materials in meltable particles and delivery through a printer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes a novel method for applying substances (I),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 38; 77pp; German.
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30-JUL-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (DEKR-) DEUT KREBSFORSCHUNGSZENTRUM. (EUMO-) EURO LAB MOLEKULARBIOLOGIE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-DEC-1999;
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                                                              Similarity
8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Breitling F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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99DE-01035553.
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                                                                                             100.0%; Score 47; DB 3; 100.0%; Pred. No. 2.1e+06
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                                                              0,
                                                              Mismatches
                                                                                          2.1e+06;
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                                                                                                                         Length 8;
                                                              Indels
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                                                          Gaps
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RESULT 94
AAB22877
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                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 8; Conserv
AAB22877;
                              AAB22877 standard; peptide; 8
                                                                                                                                                                                                                                                                        Sequence 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRO201; human; NSP; SH2-containing protein; cytostatic; modulator; Src Homology-2 containing protein; Nsp1; PRO308; Nsp2; PRO309; Nsp2 cellular response; tumorigenesis; integrin stimulation; signalling.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New isolated nucleic acid molecules encoding novel Src Homology-2 containing proteins capable of modulating the activity of cellula:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 3; Col 71-72; 63pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 response to external signaling.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Stewart TA;
                                                                                                                                                             рукрарарк в
                                                                                                                                                                                             100.0%; Score 47; DB 3; ilarity 100.0%; Pred. No. 2.1e+06; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99US-00369028
                                                                                                                                                                                                                                Length 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      of cellular
                                                                                                                                                                                             ٥,
                                                                                                                                                                                             Gaps
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10-JAN-2001

(first entry)

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activity. Such protease biodetector flusion proteins comprise one or more fluorescent proteins; a recognition signal which is cleaved by the protease; and at least one cellular localisation signal. The latter two components may be components of a single protein which is acted upon by the protease, or may be from heterologous sources. Due to the localisation signal, the biodetector protein is localised to a particular region of the cell. Once acted on by the protease of interest, the fluorescent protein is cleaved from the localisation sequence, and is free to migrate to other locations within the cell. The presence of a second localisation signal attached to the fluorescent protein enables the fluorescent protein to be directed to a different cellular compartment after cleavage of the protease recognition sequence. The compartment after cleavage of the protease recognition. The methods and biosensors of the invention can be used to investigate a wide range of cellular activities and to screen compounds which modulate these cativities. Biosensors containing a recognition site for caspase, for example, may be used for the screening of compounds which modulate these capoptosis, while biosensors containing other protease recognition sites anthrax
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-FEB-1999;
08-MAR-1999;
12-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            drug
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to systems, methods and reagents for cell-based screening or detection of compounds which affect particular biological functions. The methods of the invention utilise fluorescent biodetector molecules which, when acted on by a compound of interest, cause an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Automated cell-based characterization of toxin by contacting cells containing luminescent reporter molecules with test substance and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Giuliano KA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-FEB-2000; 2000WO-US004794.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              31-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Biodetector protein; fusion protein; recognition site; cellular targetting sequence; cellular localisation; fluorescent protein protease activity detection; toxin detection; cellular stress detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FLAG epitope,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 alteration in the cellular distribution of at least the fluorescent moiety. In one embodiment, the biosensors comprise heat shock proteins (HSPs) fused to a fluorescent protein (e.g., jellyfish green fluorescent protein (GFP), or derivatives thereof). Such biosensors are located in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 11; Fig 29A; 336pp; English.
                                                                                 candidate compound optimisation by combining many cell screening form with fluorescence-based molecular reagents and computer-based feature extraction, data analysis and automation, resulting in increased quant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the cytoplasm, but on stress activation translocate to the nucleus. In another embodiment biodetector proteins can be used to detect protease
                                                                                                                                                     may be used for the detection of proteolytic toxins (such as anthrax lethal factor). The method provides improved target validation and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CELL-) CELLOMICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              discovery;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         optically.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO:36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99US-0123399P.
99US-00352171.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99US-0122152P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cell based screening; epitope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        fluorescent protein;
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and speed of data collection and faster evaluation of drug candidates Sequences AAB22877-B22880 represent epitope sequences that can be use biosensor fusion proteins of the invention

quantity formats

AAB28432 ID AAB: XX

AAB28432 standard; peptide; 8 AA

RESULT

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ARBSULT 95
AAB26126
AC AAB26
XX Humar
XX Humar
XX PCIGE
KW PCIGE
K
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                                                                                                                                         Best
                                                                                                              Matches
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                                                                                                                                                                                                                                                                                   The present sequence comprises the translated sequence of an affinity purification recognition system site. It was inserted into a vector for the human prostate cancer-induced growth factor (PCIGF). The protein is member of the TGF-beta superfamily. The 3' part of its gene has been shown to be upregulated in prostate cancer, and thus can be used in the diagnosis of the disease. It is also useful for the identification of antagonists to the PCIGF protein which can be used to treat cancer. In addition, benign prostatic hyperplasia, prostatitis and prostatic intraepithelial neoplasia can be detected using the gene and protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel prostate cancer induced growth factor derived polynucleotide useful for detecting, diagnosing, prognosing, preventing or treating conditions of the prostate, especially prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Billing-Medel
Russell JC, (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; TGF-beta superfamily; prostate cancer-induced growth factor; PCIGF; prostate disease; prostate cancer; benign prostatic hyperpla prostatitis; prostatic intraepithelial neoplasia; affinity purifica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human TGF-beta superfamily protein PCIGF recognition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAB26126 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-628218/60.
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                                                                                                                                                                                                                             Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 11a; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ABBO ) ABBOTT LAB
                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 DYKDDDDK 8
                                      1 DYKDDDDK 8
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8; Conser
                                                                                                              Similarity
8; Conserv
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                                                                                                              Conservative
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                                                                                                                                            100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            117pp;
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Pred. No. 2.1e+06;
                                                                                                                                            Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                 Mismatches
                                                                                                                                            2.1e+06;
                                                                                                                                                                             DB 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gordon
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                                                                                                                                                                       Length 8;
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RESULT 97
AAY95914
ID AAY95
XX AAY95
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DT 20-NC
XX FLAG
XX FLAG
XX APOPT
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Matches
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                                                                                                                                                               FLAG; Goodpasture antigen binding protein; GPBP; apoptosis; cancer; tumour; diagnosis; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               an artificial mini-exon used in a novel gene-trapping retrovirus. The retrovirus comprises a branch point sequence, a polypyrimidine tract, a splice acceptor, a splice donor and long-terminal repeats (LTRs). It is useful for introducing a mutation into a gene in a cell, for determining the expression pattern of a gene in a non-human animal, and for gene therapy. It is also useful for producing transgenic animals such as mice, zebrafish, pufferfish, medaka, frogs, flies, goats, sheep, cows, pigs and chickens. High-titre virus producer cell lines have been developed which permit efficient production of retroviral vectors for the construction of
                                                                                                                                                                                                                                           FLAG peptide
                                                                                                                                                                                                                                                                                                20-NOV-2000
                                                                                                                                                                                                                                                                                                                                                    AAY95914;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 8 AA;
                                                                      WO200050607-A2
                                                                                                                                                                                                                                                                                                                                                                                                 AAY95914 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polypyrimidine tract,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-628260/60.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 DYKDDDDK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            рукрараж в
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      t retrovirus for gene trapping, gene mutation and for animal production, comprises a branch point sequence, dine tract, splice acceptor, splice donor and long-terminal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  animals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chen
                                                                                                                                                                                                                                                                                                (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gene
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                                                                                                                                                                                                                                                                                                                                                                                              peptide; 8
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                                                                                                                                                                                                                                                                                                entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the FLAG epitope,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.1e+06;
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                                                                                                                                                                                            disease;
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RESULT 98
AAY53260
Query Match
Best Local Similarity
""" hes 8; Conserva
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                                                                                              07-MAY-1996;
05-MAY-1997;
16-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence is that of FLAG peptide. FLAG was expressed at the N -terminus of recombinant human Goodpasture antigen binding protein (GPBP) constructs (see AAY95907-11) of the invention to facilitate purification from host cells. GPBP (see AAY95900) is a novel serine/threonine kinase that specifically binds to and phosphorylates the N-terminus of Goodpasture antigen. The invention provides nucleic acids (see AAA50341-53) encoding GPBP, recombinant vectors, host cells, encoded polypeptides (see AAY95900-11) and antibodies. It also provides methods for detecting the presence of an autoimmune condition or apoptosis by detecting an increase in GPBP expression, and methods for treating an autoimmune
                                                                                                                                                                                                                                                                                                 antiinflammatory; antiarteriosclerotic; chemotaxis inhibitor; brain;
viral encephalitis; multiple sclerosis; meningitis; head trauma; stroke;
neuro-degenerative disease; HIV encephalopathy; primary brain tumour;
                                                                                                                                                                                     28-MAR-2000
                                                                                                                                                                                                                                             Mus musculus.
                                                                                                                                                                                                                                                                                                                                             Neurotactin; identification; characterisation; chemokine; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                         17-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY53260 standard; peptide; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 8
                                                                                                                                                                                                                                                                         respiratory intection;
                                                                                                                                                                                                                                                                                                                                                                            Mouse neurotactin
                                                                                                                                                                                                                                                                                                                                                                                                                                      AAY53260;
                                                                   (MILL-)
                                                                                                                                                                                                               JS6043086-A
                                                                                                                                                                                                                                                                                       lupus associated cerebritis; inflammatory disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel Goodpasture antigen binding treating autoimmune disorders, tun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-572094/53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 22; 158pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-FEB-2000; 2000WO-IB000324
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                                                                 MILLENIUM BIOTHERAPEUTICS
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97US-00851160.
97US-00991426.
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                                                                                                                                                                                                                                                                         diagnosis;
                                                                                                                                                                                                                                                                                                                                                                          epitope tag SEQ ID NO:14.
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                                                                                                                                                                                                                                                                       gene therapy.
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WPI; 2000-270344/23

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ID AAX9
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KW Cel:
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Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              respiratory infections. Anti-neurotactin antibodies are useful in diagnostic assays for monitoring the level of neurotactin produced by a mammal and in conjunction with gene therapy they are useful to evaluate the normal and engineered neurotactin-expressing cells prior to their introduction into the patient. (I) are useful as labels or targets for killing cells which express cell surface neurotactin and for purification of natural or recombinant neurotactin from cells. (I) have a longer half-life within the human body than other antibodies. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        lymphocytes by blocking binding of neurotactin to its receptor. (I) are useful for treating disorders-associated with abnormal expression or activity of neurotactin which includes inflammation of the brain associated with disorders such as viral encephalitis, multiple sclerosis, viral or bacterial meningitis, head trauma, stroke, neuro-degenerative diseases, HIV encephalopathy, primary brain tumours and lupus associated cerebritis and also for inflammatory diseases e.g. atherosclerosis or
                                                                                                                                                                                                                                                                                                                                                                                            Thymic stromal lymphopoietin; TSLP; B cell maturation; immune regulation; cell proliferation; cell differentiation; cell death; cell migration; cell-to-cell interaction; inflammatory response; chromosome 5g21-22; Gardner syndrome; adenomatous polyposis coli; hereditary desmoid disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FLAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel antibodies raised against membrane-anchored chemokine, neurotactin useful for diagnosis and treatment of disorders associated with abnormal expression of neurotactin, especially inflammation of brain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY97366 standard;
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New human thymic stromal lymphopoietin (TSLP) polypeptide useful stimulating lymphocyte development and proliferation.
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                                                           WPI; 2000-387794/33.
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8; Conserv
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Pred. No. 2.1
D; Mismatches
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hes 0;
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Best Local S
Matches 8
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            Compounds that modulate the activity of a multiple lineage kinase protein (MLK) and promote either cell survival or cell death can be identified by treating a cell that contains MLK with a test compound and determining it decreases or increases the activity of MLK and promotes cell survival or death. Compounds identified as having MLK modulating activity have applications as anti-neurodegenerative agents, antiinflammatory agents and anticancer agents and are potentially useful for treatment of neurodegenerative diseases (e.g. Alzheimer's, Huntington's and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Multiple linkage kinase; MLK; PYK; modulation; antiinflammatory; anticancer; Alzheimer's disease; Huntington's disease; Parkinson's disease; amyotrophic lateral sclerosis; ischaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAY83142;
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                                                                                                                                                                                            Identifying compounds that modulate multiple lineage kinase useful e.g. for treating neurodegeneration or cancer, from ton survival or death of kinase-expressing cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                  09-MAR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200013015-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FLAG epitope.
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                                                                                                                                                               Example 14; Page 51; 158pp;
                                                                                                                                                                                                                                                                                                 Maroney A,
                                                                                                                                                                                                                                                                                                                                                                 26-AUG-1998;
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MOL. Gen. Genet. 233, 217-224, 1992
MOL. Gen. Genet. 233, 217-224, 1992
A;Title: Methanococcus voltae harbors four
A;Reference number: A59304; MUID:92293118;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Molecule type: DNA
A; Residues: 1-99, 'TA', 102-107, 'P', 126, 'LH', 129-306 <HAL>
A; Residues: 1-99, 'TA', 102-107, 'P', 126, 'LH', 129-306 <HAL>
A; Cross-references: UNIPARC: UPI0000174FE0; EMBL, X61203
A; Note: the sequence is revised in GenBank entry X61203,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Cross-references: UNIPROT:Q00406; UNIPARC:UPI00000625CA; R;Halboth, S.; Klein, A. submitted to the EMBL Data Library, August 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               methylviologen-reducing hydrogenase chain vhcG [similarity] -
C;Species: Methanococcus voltae
C;Date: 18-Aug-2000 #sequence_revision 18-Aug-2000 #text_chang
C;Accession: S32834; S16726
  R;Parkhill, J.; Wren, B.W.; Thomso deno-Tarraga, A.M.; Chillingworth,
                                                                                                                        AC0184
                                                                                                                                          RESULT 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Reference number: $16721
A; Accession: $16726
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A; Residues: 1-423 < MUR>
A; Cross-references: UNITROT: 004921;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  submitted to the EMBL Data Library, August 1991 A;Description: Methanococcus voltae harbors two
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Reference number: A; Accession: S32834
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                                                                                                                                                                                                                                                                                                                                                     Superfamily: methyl viologen-reducing
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B.W.; Thomson, N.R.; Titball, R.W.; Holden, M.T.G.; P. hillingworth, T.; Cronin, A.; Davies, R.M.; Davis, P.;
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87.5%;
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                                                                                                                                                                                                                                                                                              Score 38;
Pred. No.
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PMID:1603063
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                                                      #text_change 05-Oct-2004
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                                                                                                 (EC 1.1.1.38)
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  Prentice,
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                                                      C;Accession: S54451; S52836
R;Gentles, S.; Bowman, S.
submitted to the EMBL Data Library, May 1995
                                                                                                           hypothetical protein YMR076c - yeast (Saccharomyces N;Alternate names: hypothetical protein YM9582.01c; C;Species: Saccharomyces cerevisiae C;Date: 08-Jul-1995 #sequence_revision 09-Mar-1996 #
                    A; Reference number: S54451
A; Accession: S54451
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type:

DNZ

#text_change 09-Jul-2004

cerevisiae)

hypothetical

protein YM9916.15c

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C.Access.....P.
R;Cherel, I.; Thuriaux, P.
Yeast 11, 261-270, 1995
A;Title: The IFH1 gene product interacts with a fork head
A;Title: momber: S55352; MUID:95304839; PMID:7785326
RESULT 6
S54451
                                                                                                                                                                                                                                                                                                                                    submitted to the EMBL Data Library, A;Description: The sequence of S. c.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Molecule type: DNA
A;Residues: 1-565 <KUR>
A;Cross-references: UNIPROT:Q8ZG09; UNIPARC:UPI00000DCE2B; GB:AL590842; PIDN:CAC90334.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A,Title: Genome sequence of Yersinia pestis, A;Reference number: AB0001; MUID:21470413; PA;Accession: AC0184
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                                                                                                                                                                                                                                                                                                                                                                     R;Hallsworth,
                                                                                                                                                                                                                                                                                                                                                                                                   A; Molecule type: DNA
A; Residues: 1-1085 < CHE>
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C;Accession: S55352; S51446; S47477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IFH1 protein - yeast (Saccharomyces cerevisiae)
N;Alternate names: protein L8083.9; protein YLR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    il, M.; Rutherford, K.; S. Nature 413, 523-527, 2001
                                                                                                                                                                                                                                                                      A;Molecule type: DNA
A;Residues: 1-1085 <HAL>
                                                                                                                                                                                                                                                                                                        A;Accession:
                                                                                                                                                                                                                                                                                                                     A;Reference number: S51443
                                                                                                                                                                                                                                                                                                                                                                                    A; Cross-references: UNIPROT: P39520;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C; Species: Saccharomyces cerevisiae
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Best Local
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;Keywords: oxidoreductase
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85.7%;
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Pred. No. 1.2e
1; Mismatches
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Pred. No. 62;
2; Mismatches
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1.2e+02;
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probable spindle pole body component bimB - Emericella nidulans N,Alternate names: cell division-associated protein bimB C,Species: Emericella nidulans. Aspergillus nidulans C;Date: 10-Jun-1993 #sequence_revision 18-Nov-1994 #text_change C;Accession: A42854; S27411
R;May, G.S.; McGoldrick, C.A.; Holt, C.L.; Denison, S.H. J, Biol. Chem. 267, 15737-15743, 1992
A;Title: The bimB3 mutation of Aspergillus nidulans uncouples DN A;Reference number: A42854; MUID:92348436; PMID:1639810
A;Accession: A42854
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A42854
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A;Cross-references: UNIPROT:Q04264; UNIPARC:UPI0000168AB1; EMBL:Z49259; NID:g807956; A;Experimental source: strain AB972
R;Pearson, D.; Bowman, S.
submitted to the EMBL Data Library, April 1995
A;Reference number: S52814
                                                                             A;Note: sequence extracted from NCBI backbone (NCBIN:112444, NCBIP:112445) R;May, G.S.; McGoldrick, C.A.; Denison, S.H.; Holt, C.L. submitted to the EMBL Data Library, February 1992 A;Description: The bimB3 mutation of Aspergillus nidulans uncouples DNA re
submitted to the EMBL Data Library, A;Description: Cloning and sequenci A;Reference number: Z17060 A;Accession: T10503
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C; Keywords: nucleus
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A;Residues: 1-2067 <MAY>
A;Cross-references: UNIPROT:P33144; UNIPARC:UPI000012693F
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A;Residues: 659-1277 <PEA>
A;Cross-references: UNIPARC:UPI0000168AC7; EMBL:Z48952; NID:g763008; PIDN:CAA88801.1;
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                                                                                                                                                                                                    T10503
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;Residues: 1-2010, 'PLSIG',2016 <MA2>
;Cross-references: UNIPARC:UP100001688E9, EMBL:M83232; NID:g168020; PID:g168021;
;Note: this sequence differs by a frameshift after residue 2010
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Best Local
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Map position: 13R
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larity 85.7%;
Conservative
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                                           sequencing of
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Pred. No. 1.5e+02;
1; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                  Score 38; DB 2;
Pred. No. 2.4e+02;
                                                                    January
                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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    A;Status: preliminary
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R;Nolling, J.; Breton, G.; Omelchenko, M.V.; Markarova, K.; Daly, M.J.; Bennett, G.N.; Koonin, E.V.; Smith, D.R. J. Bacteriol. 183, 4823-4838, 2001
A;Title: Genome Sequence and Comparative Analysis of the A;Reference number: A96900; MUID:21359325; PMID:21359325 A;Accession: B96974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khaykin, E.; C.A.; Li, J.H.; Li, Y.; Lin, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Maiti, R.; Martizzo, M.; Rooney, T.; Rowley, D.; Sakano, H. A;Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sun, H.; Taker, M.; Wu, D.; Yu, G.; Fraser, C.M.; Venter, J.C.; Davis, R.W. A;Title: Sequence and analysis of chromosome 1 of the plant Arabidopsis. A;Reference number: A86141; MUID:21016719; PMID:11130712
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A;Molecule type: mRNA
A;Residues: 1-351 <MUT>
A;Cross-references: UNIPROT:023791; UNIPARC:UPI00000A4EB5;
A;Experimental source: cv. N67-10; nearly mature fruit
C;Function:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chin, C.W.; Chung, M.K.; Conn, L.; ansen, N.F.; Hughes, B.; Huizar, L. Nature 408, 816-820, 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             F;25-121/Domain: amino-terminal propeptide #status predicted <PRO>
F;122-339/Product: fruit bromelain #status predicted <MAT>
F;340-351/Domain: carboxyl-terminal propeptide #status predicted <CTP>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hypothetical protein F11A17.5 [imported] - Arabidopsis thaliana
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 09-Jul-2004
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                                                                                                                                                C;Date: 14-Sep-2001 #sequence_revision 14-Sep-2001 #text_change 09-Jul-2004 C;Accession: B96974
                                                                                                                                                                                                  superfamily I DNA and RNA helicase [imported] - Clostridium acetobutylicum C_iSpecies: Clostridium acetobutylicum
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C; Superfamily:
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A; Residues: 1-513 <STO>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Accession: B96524
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Arabidopsis thaliana hypothetical protein F17J16.30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       UNIPROT:Q9SX72; UNIPARC:UPI00000A3573; GB:AE005173; NID:g5733869; P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               78.7%;
85.7%;
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85.7%;
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                                                                                                                              K.s.;
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                                                                                                                              Q.; Gibson,
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RESULT 11
T20815
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A;Residues: 1-721 <KUR>
A;Cross-references: UNIPROT:Q97LF8; UNIPARC:UPI00000C9F00; GB:AE001437; PIDN:AAK78581.1;
A;Experimental source: Clostridium acetobutylicum ATCC824
C;Genetics:
     A; Map position: 4
A; Introns: 86/1; :
A; Note: F10M6.180
                                                         A;Cross-references: UNIPARC:UPI000016DADA;
A;Experimental source: cultivar Columbia; E
C;Genetics:
                                                                                                                                                                                                                               hypothetical protein F10M6.180 - Arabidopsis thaliana
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 23-Apr-1999 #sequence_revision 23-Apr-1999 #text_change 23-Jul-1999
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C;Superfamily: Bacillus subtilis hypothetical protein yvgS
                                                                                                                                               A; Reference number: Z15414
A; Accession: T05410
                                                                                                                                                                                                                                                                                                              RESULT 12
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A; Residues: 1-791 <WI2>
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A;Experimental source: clone F13A7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: DNA
A; Residues: 1-791 <WIL>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     submitted to the EMBL Data Library, A;Reference number: Z19327
A;Accession: T20815
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A;Accession: T21409
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                                                                                                          ;Molecule type: DNA;Residues: 1-838 <BEV>
                                                                                                                                                                                              ;Accession: T05410 ;Bevan, M.; Weichselgartner, M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ;Map position: 5
;Introns: 28/3; 194/2; 195/3; 299/2; 404/1; 462/3; 500/1; 670/3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cross-references: UNIPARC:UPI0000060F93; Experimental source: clone F26D2
                                                       Genetics:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Species: Caenorhabditis elegans
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les 6; Conserv
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                      158/1; 175/3;
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85.7%;
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75.0%;
                      214/3;
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Pred. No. 1.3e
0; Mismatches
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Database,
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Pred. No.
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                    254/3; 286/2; 317/3; 361/3;
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                                                                          BAC
                                                                        EMBL:AL021811
BAC clone F10M6
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1.2e+02;
                                                                                                                                                                               B.; Granderath,
February 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      .3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 791;
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                    399/2; 463/3; 533/2;
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                                                                                                                                                                                                D.; Herzl,
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                                                                                                                                                                                                 A.;
RESULT 15
S39814
DNA polymerase homolog
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R;Gattung, S.; Scheet, P.
submitted to the EMBL Data Library, July 1997
submitted to the EMBL Data Library, July 1997
A;Description: The sequence of C. elegans cosmid C33C12.
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                                                                                                                                                                                                                                                                                                                                                                                                         hypothetical protein C33C12.7 - Caenorhabditis elegans
C;Species: Caenorhabditis elegans
C;Date: 29-Oct-1999 #sequence_revision 29-Oct-1999 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Experimental source: strain B31 C;Superfamily: Borrelia burgdorferi hypothetical protein BB0739
                                                                                                                                         A; Introns: 63/3; 94/3; 124/2; 168/2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nature 390, 580-586, 1997
A;Authors: Smith, H.O.; Venter, J.C.
A;Title: Genomic sequence of a Lyme disease
A;Reference number: A70100; MUID:98065943; P
A;Accession: B70192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Fraser, C.M.; Casjens, S.; Huang, W.M.; Sutton, G.G.; Clayton, R.; Lathigra, R.; son, D.; Peterson, J.; Kerlavage, A.R.; Quackenbush, J.; Salzberg, S.; Hanson, M.; Bowman, C.; Garland, S.; Fujii, C.; Cotton, M.D.; Horst, K.; Roberts, K.; Hatch,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT
B70192
                                                                                                                                                          A; Map position:
                                                                                                                                                                            A;Gene: CESP:C33C12.7
                                                                                                                                                                                                               A;Cross-references: UNIPROT:016579; UNIPARC:UPI0000078124; EMBL:AF016662; PIDN:AAB66055
A;Experimental source: strain Bristol N2; clone C33C12
                                                                                                                                                                                                                                               A; Molecule type: DNA
A; Residues: 1-215 <GAT>
                                                                                                                                                                                                                                                                                                   A; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A_iStatus: preliminary; nucleic acid sequence not shown; translation not A_iMolecule type: DNA
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                                                                                                                                                                                                                                                                                                                      A;Reference number: Z21105
                                                                                                                                                                                                                                                                                                                                                                                            C;Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: UNIPROT:OS1681; UNIPARC:UPI00000575C8; GB:AE001174; GB:AE000783;
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                                                                                     Score 36;
Pred. No.
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Pred. No.
                                                                      Mismatches
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PMID:9403685
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49;
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pol-r -

rye mitochondrion (strain Halo)

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A; Variety:
C; Date: 19-
C; Accession
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A; Variety:
C; Date: 13-
C; Accession
                                                                 hypothetical protein F7K15.80 - Arabidopsis thaliana (,Species: Arabidopsis thaliana (mouse-ear cress) C;Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 02-Jun-2000 #sequence_revision 02-Jun-2000 #text_change 09-Jul-2004 C;Accession: T49237
R;Obermaler, B; Ottenwaelder, B.; Duchemin, D.; Zeitler, K.; Mewes, H.W.; submitted to the Protein Sequence Database, April 2000 A;Reference number: Z25019
A;Accession: T49237
                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Residues: 1-312 <DOH>
A;Cross-references: UNIPROT:Q36546;
A;Experimental source: strain B-P2;
A;Note: the nucleotide sequence was
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                              A;Gene: pol-r
A;Genome: mitochondrion
C;Keywords: mitochondrion
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R;Dohmen, G.; Tuu
Genet. 25,
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S39823
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Renet. 25,
A;Residues: 1-485 <OBE>
A;Cross-references: UNIPROT:Q9LXL1; UNIPARC:UPI00000A1EFC; EMBL:AL353871; GSPDB:GN00061;
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A;Status: nucleic acid sequence
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                                     A; Molecule type: DNA
                                                   A; Status: preliminary
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;Cross-references: UNIPROT:Q36545;
;Experimental source: strain Halo;
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19-Mar-1997 #sequence_revision
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25, 59~65, 1994
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25, 59-65, 1994
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Pred. No. 72;
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Pred. No. 67;
                                                                                                                      Duchemin, D.; Zeitler, K.; Mewes, H.W.; Rudd,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              UNIPARC:UPI0000096B6D; EMBL:X74133; NID:g4392 etiolated shoots submitted to the EMBL Data Library, July 1993
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R;Lin, X.; Kaul, S.; Rounsley, S.D.; Shea, T.P.; Benito, M.I.; Town, C.D.; Fujii, C.Y.; M.; Koo, H.; Moffat, K.S.; Cronin, L.A.; Shen, M.; VanAken, S.E.; Umayam, L.; Tallon, Leuss, D.; Nierman, W.C.; White, O.; Eisen, J.A.; Salzberg, S.L.; Fraser, C.M.; Venter,
                                                                                                                                                                                                                                                                                                                                            hypothetical protein YFL023w - yeast (Saccharomyces cerevisiae) C;Species: Saccharomyces cerevisiae C;Date: 02-Sep-1995 #sequence_revision 19-Oct-1995 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            probable ATP-dependent RNA helicase [imported] - Arabidopsis thaliana
(;Species: Arabidopsis thaliana (mouse-ear cress)
C;Dace: 02-Feb-2001 #sequence_revision 02-Feb-2001 #text_change 09-Jul-2004
C:Accession: H84913
                                                                                                   A; Cross-references:
A; Map position: 6L
                                                                                                                                                                                       A; Molecule type: DNA
A; Residues: 1-796 < MUR>
                                                                                                                                                                                                                                                                            submitted to the EMBL Data A; Description: Analysis of
                                                                                                                                                                                                                                                                                                               R; Murakami, Y.; Naitou, M.; Hagiwara, H.;
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A; Residues: 1-748 <STO>
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                                                                                                                                                                        A;Cross-references: UNIPROT:P43573;
                                                                                                                                                                                                                                                            A;Reference number: S56186
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                                                                 Length 796;
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R;anonymous, The C. elegans C. elegans: a platform Science 282, 2012-2018, 1998
Science 282, 2012-2018, 1998
A;Title: Genome sequence of the nematode C. elegans: a platform A;Title: Genome sequence MUID:99069613; PMID:9851916
A;Note: see websites genome wustl.edu/gsc/C_elegans/ and www sar 
A; Description: The sequence
A; Reference number: $50432
A; Accession: $50536
A; Molecule type: DNA
A; Residues: 1-1076 <DIE>
                                                                                                                                    hypothetical protein YER033c - yeast (Saccharomyces cerevisiae) C;Species: Saccharomyces cerevisiae C;Date: 28-May-1993 #sequence_revision 24-Feb-1995 #text_change C;Accession: S50536 R;Dietrich, F.S. submitted to the EMBL Data Library, December 1994
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A88746
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C;Species: Caenorhabditis elegans
C;Date: 10-May-2001 #sequence_revision 10-May-2001:
C;Accession: A88746
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A;Reference number: S07183
A;Accession: S07183
A;Status: translation not sho
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C; Superfamily: DNA polymerase,
C; Keywords: mitochondrion
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R;Paillard, M.; Sederoff, R.R.; Levings III, C.
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C;Genetics:
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A; Residues: 1-917 < PAI>
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probable membrane associated protein PFB0190c - malaria parasite (Plasmodium C;Species: Plasmodium falciparum C;Date: 13.Nov-1998 #sequence_revision 13-Nov-1998 #text_change 09-Jul-2004 C;Date: 13.Nov-1998 #sequence_revision 13-Nov-1998 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Residues: 1-1287 < DUZ>
A; Cross-references: UNIPROT:Q18100; UNIPARC:UPI0000081960; EMBL:U50067;
A; Cross-references: UNIPROT:Q18100; UNIPARC:UPI0000081960; EMBL:U50067;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R.Du, Z.; Gattung, s.
submitted to the EMBL Data Library, August
submitted to the EMBL Data Library, August
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                                                                                                                                                      A; Molecule type: DNA
A; Residues: 1-2295 <
                                                                                                                                                                                                       A;Title: Chromosome 2 sequence of the human malaria parasite Plasmodium falciparum A;Reference number: A71600; MUID:99021743; PMID:9804551 A;Accession: B71621
                                                                                                                                                                                                                                                      R;Gardner, M.J.; Tettelin, H.; Caru.; Pertea, M.; Salzberg, S.; Zhou, Science 282, 1126-1132, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Date: 22-Oct-1999 #sequence_revision C;Accession: T30988
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C;Genetics:
A;Gene: SGD:ZRGB; MIPS:YER033C
A;Cross-references: SGD:S0000835
                                                                                                                                                                                      A; Status: preliminary; nucleic acid sequence
                                                                                                                                                                                                                                                                                                        C; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Note: C18F3.2
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A; Introns: 34/3; 7
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                              Query Match
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Matches 6
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Caenorhabditis elegans
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ilarity 100.0%;
Conservative
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85.7%;
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                              k; Score 36; DB
k; Pred. No. 5.9
0; Mismatches
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Pred. No. 3.2e
1; Mismatches
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2.7e+02;
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                                              .9e+02
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R.; White, O
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C;Species:
C;Date: 15
C;Accessic
R;Lawson,
                                heat shock protein hsp18 - Clostridium acetobutylicum (Species: Clostridium acetobutylicum C;Date: 07-Apr-1994 #sequence revision 07-Apr-1994 #text C;Accession: A40592; 825534 R;Sauer, U.; Duerre, P.

J. Bacteriol. 175, 3394-3400, 1993 J. Bacteriol. 175, 3394-3400, 1993 A;Title: Sequence and molecular characterization of a Dh A;Reference number: A40592; MUID:93273706; PMID:8501044 A;Accession: A40592
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: UNIPROT:Q8YP82; UNIPARC:UPI00000CEAB9; GB:BA000019; A;Experimental source: strain PCC 7120 C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Note: Nostoc sp. strain PCC 7120 is a synonym of Anabaena sp. strain PCC 7120 (;Date: 14-Dec-2001 #sequence_revision 14-Dec-2001 #text_change 09-Jul-2004 C;Accession: AF2345 R;Kaneko, T.; Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, A.; Irigu Nakazaki, N.; Shimpo, S.; Sugimoto, M.; Takazawa, M.; Yamada, M.; Yasuda, M.; Tabata DNA Res. 8, 205-213, 2001 A;Title: Complete Genomic Sequence of the Filamentous Nitrogen-fixing Cyanobacterium A;Reference number: AB1807; MUID:21395285; PMID:11759840 A;Accession: AF2345
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Status: preliminary; translated
A;Molecule type: DNA
A;Residues: 1-2457 <LAW>
A;Cross-references: UNIPROT:077375
C;Genetics:
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submitted to the EMBL Data Library,
A; Reference number: 218935
A; Accession: T18492
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A; Residues: 1-78 < KUR>
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C;Superfamily: Plasmodium
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C;Species: Plasmodium falciparum
C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 09-Jul-2004
A;Molecule
                                                                                                                                                                                                                                            A40592
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                Status: preliminary
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6; Conserv
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85.7%;
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Pred. No. 6.4e+02;
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Pred. No.
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                                                                            DNA region encoding a small heat
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morecular chaperone (small heat shock protein), HSP18 [imported] - Clostridium acetobut C;Species: Clostridium acetobutylicum C;Date: 14-Sep-2001 #sequence_revision 14-Sep-2001 #text_change 09-Jul-2004 C;Accession: G97355
                                                                                                                                                                                                                                                                                                                                     hypothetical protein C03H5.3 - Caenorhabditis elegans
C;Species: Caenorhabditis elegans
C;Date: 29-Oct-1999 #sequence_revision 29-Oct-1999 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Residues: 1-151 <KUR>
A;Cross-references: UNIPROT:Q03928; UNIPARC:UPI000012CC00; GB:AE001437; PIDN:AAK81634.1
A;Experimental source: Clostridium acetobutylicum ATCC824
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G97355
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R;Rohlfing, T.; Wohldmann,
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C;Superfamily: alpha-crystallin-related small heat shock protein
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                                                                     A; Introns: 2/1;
                                                                                                        A; Gene: CESP: C03H5.3
                                                                                                                                          A; Experimental source:
                                                                                                                                                                             A; Molecule type: DNA
A; Residues: 1-287 < ROH>
                                                                                                                                                                                                             A;Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                    A; Accession: T32029
                                                                                                                                                                                                                                                   A;Description: The sequence of A;Reference number: Z21114
                                                                                                                                                                                                                                                                   submitted to the EMBL Data Library, July 1997 A;Description: The sequence of C. elegans cosmid C03H5
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                  Best
                                   Query Match
                                                                                                                                                              Cross-references: UNIPROT:016657;
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Similarity 5; Conserv
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 Conservative
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                                                                                                                                            strain Bristol
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Score 35; DB
Pred. No. 98;
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unknown protein, 9003-8083 [imported] - Arabidopsis thaliana
C.Species: Arabidopsis thaliana (mouse-ear cress)
C.Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 09-Jul-2004
C.Accession: D96708
R.Theologis, A.; Ecker, J.R.; Palm, C.J.; Federspiel, N.A.; Kaul, S.; White, O
Chin, C.W.; Chung, M.K.; Conn, L.; Conway, A.B.; Conway, A.R.; Creasy, T.H.; I
ansen, N.F.; Hughes, B.; Huizar, L.
Nature 408, 816-820, 2000
A;Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khaykin, E
C.A.; Li, J.H.; Lin, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Waiti, R.;
A;Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sun, H.
Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.
R;Title: Sequence and analysis of chromosome 1 of the plant Arabidopsis.
A;Reference number: A86141; MUID:21016719; PMID:11130712
RESULT 32
PC4365
calcium-binding protein - tapeworm
C;Species: Echinococcus granulosus
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A;Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sun, H.; Tallon, ker, M.; Wu, D.; Yu, G.; Fraser, C.M.; Venter, J.C.; Davis, R.W.
A;Title: Sequence and analysis of chromosome 1 of the plant Arabidopsis.
A;Reference number: A86141; MUID:21016719; pMID:11130712
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A;Molecule type: DNA
A;Residues: 1-306 <STC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    unknown protein [imported] - Arabidopsis thaliana C;Species: Arabidopsis thaliana (mouse-ear cress) C;Space: 02-Mar-2001 #sequence_revision 02-Mar-200
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;Accession: D96708
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85.7%;
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Pred. No. 1e+C
O; Mismatches
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Maiti, R.; Ma
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Maiti, R.; Marziali
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war, K.;
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  Query Match
Best Local Similarity
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A;Status: preliminary; translated from GB/EMBL/DDBJ A;Molecule type: DNA A;Molecule type: DNA A;Residues: 1-410 <WILL A;Residues: 1-410 <WILL A;Residues: 1-410 <WILL DROT: 002327; UNIPARC: UPI0000 A;Cross-references: UNIPROT: 002327; UNIPARC: UPI0000
                                                                                                                                                                                                                                                                                                             hypothetical protein T23D8.3 - Caenorhabditis elegans
C;Species: Caenorhabditis elegans
C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gibberellin 20-oxidase (EC 1.14.11.-) - spinach
C;Species: Spinacia oleracea (spinach)
C;Date: 11-Jun-1999 #sequence_revision 11-Jun-1999
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R;Rodrigues, J.J.S.; Ferreira, H.B.; Farias, S.E.; Zaha, A.
Biochem. Biophys. Res. Commun. 237, 451-456, 1997
A;Title: A protein with a novel calcium-binding domain associated A;Reference number: PC4365; MUID:97415820; PMID:9268732
A;Accession: PC4365
                                                                                                                                                                                                    submitted to the EMBL Data A;Reference number: Z19989 A;Accession: T25165
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Cross-references: UNIPROT:Q24795; UNIPARC:UPI00000755B8; GB:L33817; NID:g499140; PID:C;Comment: This protein plays a role in the development of the parasite by mobilizing F;1-7,8-20,21-33,34-45,46-89,90-103,104-116,117-129,130-141,142-155,156-169,170-182,18
A; Introns: 61/3;
                   A; Map position:
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A; Residues: 1-374 < WUK>
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                                           A;Gene: CESP:T23D8.3
                                                                                        A; Experimental source:
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208/3; 247/3; 396/3
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                                                                                        UNIPROT:002327;
ce: clone T23D8
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85.7%;
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Pred. No. 1.2e
0; Mismatches
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                                                                                                             UNIPARC: UPI0000078796; EMBL: Z81128;
                                                                                                                                                                                                                                                 October 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      expression of PMID:8742334
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1.3e+02
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1.2e+02;
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                                                                                                                PIDN: CAB03401.1
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C.

74.5%;

Score Pred.

35;

1.4e+02; ĎΒ

Length 410;

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RESULT
S41720
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C;Date: 27-Oct-1995 #sequence_revision 10-Nov-1995 #text_change 15-Mar-2004
C;Accession: S58653
R;Murakami, Y.; Naitou, M.; Hagiwara, H.; Shibata, T.; Ozawa, M.; Sasanuma,
          intermediate filament - goldfish
C;Species: Carassius auratus (goldfish)
C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 09-Jul-2004
C;Accession: S41720
C;Accession: S41720
R;Glasgow, E; Druger, R.K.; Fuchs, C.; Lane, W.S.; Schechter, N.
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A; Introns: 16/2
F; 399-448/Domai:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      submitted to the EMBL Data A;Description: Analysis of A;Reference number: S56186 A;Accession: S58653
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A;Experimental source: strain S288C
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A; Accession: S67161
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R;Jauniaux, J.C.; P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hypothetical protein YOR264w - yeast (Saccharomyces cerevisiae)
N;Alternate names: hypothetical protein O5375
C;Species: Saccharomyces cerevisiae
C;Date: 12-Jul-1996 #sequence_revision 12-Jul-1996 #text_change 09-Jul-2004
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EMBO
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Best Local Similarity
Matches 5; Conserv
                                                                                                                                                                                                                                                                                                                 ;399-448/Domain: SH3 homology <SH3>
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Residues: 1-451 <MUR>
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85.7%;
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Pred. No. 1.6e+02;
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Pred. No. 1.5e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
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C;Species: Plasmodium falciparum control mataria parasite (Plasm C;Date: 13-Nov-1998 #sequence_revision 13-Nov-1998 #text_change 09-Jul-2004 C;Accession: G71615
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-472 <GLA>
                                                                                                                                                                                                                                               A;Reference number: A71600; MUID:99021743; PMID:9804551
A;Accession: G71615
                                                                                                                                                                                                                                                                                                                          R;Gardner, M.J.; Tettelin, H.
.; Pertea, M.; Salzberg, S.;
Science 282, 1126-1132, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hypothetical protein At2g47860 [imported] - Arabidopsis thaliana C;Species: Arabidopsis thaliana (mouse-ear cress) C;Species: Arabidopsis thaliana (mouse-ear cress) C;Datc: 02-Feb-2001 #sequence_revision 02-Feb-2001 #text_change 09-Jul-2004
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Query Match
Best Local Similarity
""" hes 6; Conserv:
                                                                                                                                                                                                                                                                                                                                                                                                                                    phospholipase A2-like a/b fold hydrolase PFB0410c - malaria parasite (Plasmodium C;Species: Plasmodium falciparum
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C; Superfamily: Arabidopsis hypothetical
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A; Residues: 1-635 <STO>
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                                                                                                                                                A;Cross-references: UNIPROT:096176; UNIPARC:UPI0000077A97; GB:AE001393; GB:AE001362; A;Experimental source: clone 3D7
                                                                                                                                                                                                             A; Molecule type: DNA
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Matches
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nes 6; Conserv
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                                                                                                                                                                                                1-679 <GAR>
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                                                                                                                                                                                                                                                                                                                                            Tettelin, H.; Carucci, D.J.; Cummings, L.M.; Aravind, L.; Koonin, slzberg, S.; Zhou, L.; Sutton, G.G.; Clayton, R.; White, O.; Smith,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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75.0%;
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Pred. No.
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Pred. No. 1.6e+02;
1; Mismatches 1
                                                                                                                                                                                                                                      sequence not shown; translation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  UNIPARC: UPI00000A0189; GB: AE002093; NID: g3738298;
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2.3e+02;
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                                                                                                                                                                                                                                                                                                        Plasmodium falciparum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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nayam, L.; Tallon, L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                two
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Conservative

<u>,,</u>

Mismatches

Indels

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Gaps

0

74.5%; 85.7%;

Score 35; I Pred. No. 2.

.4e+02;

Length 679

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RESULT 42
RNZQ2L
DNA-directed RNA polymerase (EC 2.7.7.6) II large chain -
C;Species: Plasmodium falciparum
C;Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_c
C;Accession: S07485
R;Li, W.B.; Bzik, D.J.; Gu, H.; Tanaka, M.;
Nucleic Acids Res. 17, 9621-9636, 1989
A;Title: An enlarged largest subunit of Place
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT
S46817
                                                                                                                                                                               문
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                                                                                                                                                                                                                                                                                                            A;Map position: 8R
C;Superfamily: uncharacterized conserved
C;Keywords: transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                        R;Favello, T.
submitted to the EMBL Data Library, June 1994
A;Description: The sequence of S. cerevisiae |
A;Reference number: S46795
A;Accession: S46817
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5
                                                                                                                                                                                                                                                                                                                                           A;Cross-references: SGD:S0001122
A;Map position: 8R
                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: UNIPROT:P38800;
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                         A; Molecule type: DNA
A; Residues: 1-1345 < FAV>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hypothetical protein YHR080c - yeast (Saccharomyces cerevisiae)
C;Species: Saccharomyces cerevisiae
C;Date: 28-Oct.1994 #sequence_revision 28-Oct-1994 #text_change
C;Accession: S46817
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Residues: 1-1010 <LAT>
A;Cross-references: UNIPARC:UPI000017BA5D; EMBL:U53332; PIDN:AAC71155.1; GSPDB:GN00020;
A;Experimental source: strain Bristol N2; clone F59G1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Reference number:
A;Accession: T34314
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   submitted to the EMBL Data Library, April A;Description: The sequence of C. elegans A;Reference number: Z21504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R;Latreille,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hypothetical protein F59G1.3 - Caenorhabditis elegans C;Species: Caenorhabditis elegans C;Date: 29-Oct-1999 #sequence_revision 29-Oct-1999 #t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      밁
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                                                                                                                                                                                                                                                             Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                           Genetics
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Best Local
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                                                                                                                                                                                                                                                                                                           transmembrane
                                                                                                                                                                                                                                            Similarity 6; Conserv
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                                                                                                                                                                                                               DYKODDD 7
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                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     113/1; 149/3; 188/2;
                                                                                                                                                                                                                                                           74.5%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     74.5%;
75.0%;
                                                                                                                                                                                                                                                                                                           protein
 subunit of Plasmodium
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Pred. No.
                                                                                                                                                                                                                                                           Score 35;
Pred. No.
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ches 1;
                            Fox,
                                                                                                                                                                                                                                                           5e+02;
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                              B.A.; Inselburg,
falciparum RNA
                                                            #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   #text_change
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                                                                                                                                                                                                                                                                         Length 1345;
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                                                                                            malaria parasite
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polymerase II
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                                                                                            (Plasmodium
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defines
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T32477
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                              A; Introns: 45/3;
                                                            A;Gene:
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                                                                                                                                                                                                                          submitted to the
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C; Species: P
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                                                                             Genetics:
Query Match
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Best Local
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Best Local :
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C;Accession: T32477
R;Chissoe, S.; Haw)
                                                                                                                                                                                                                                                              hypothetical protein F52H2.3 - Caenorhabditis elegans C;Species: Caenorhabditis elegans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Reference number: A71600; MUID:99021743; PMID:9804551
A;Accession: D71614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Gardner, M.J.; Tettelin, H.
; Pertea, M.; Salzberg, S.;
Science 282, 1126-1132, 1998
A;Title: Chromosome 2 sequenc
                                     A; Experimental source:
                                                          A; Cross-references:
                                                                            A; Residues: 1-140 < CHI>
                                                                                          A; Molecule type: DNA
                                                                                                             A; Status: preliminary; translated
                                                                                                                                                A;Description: The sequence
A:Reference number: Z21175
                                                                                                                                                                                                                                          A;Cross-references: UNIPROT:096185; UNIPARC:UPI0000079076; GB:AE001396; GB:AE001362; NIA;Experimental source: clone 3D7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Date: 13-Nov-1998 #sequence_revision 13-Nov-1998 #text_change
C;Accession: D71614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Accession: S07485
A;Molecule type: DNA
A;Residues: 1-2452 <LIW>
A;Cross-references: UNIPROT:P14248; UNIPARC:UPI00001345CB;
C;Genetics:
                                                                                                                               A; Accession: T32477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Molecule type: DN
A; Residues: 1-2573
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C;Superfamily: Plasmodium DNA-directed RNA polymerase II large chain C;Keywords: DNA binding; nucleotidyltransferase; phosphoprotein; tand F;68-84/Region: zinc finger CCHH motif
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Status: preliminary; nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 F;2247-2384/Region: 7-residue repr
F;376/Binding site: ATP/GTP (Lys)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Reference number:
CESP: F52H2.3
                                                                                                                                                                                                                                                                                                                                                                                1331 DYYDDDD 1337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1264 DYVDDDD 1270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Plasmodium falciparum
                                                                                                                                                                                                                                                                                                                                                                                                                                                      6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                     DYKODDD 7
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                                                                                                                                                                  EMBL Data Library, September 1997 ne sequence of C. elegans cosmid F52H2
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alzberg, S.; Zhou,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA
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                                   UNIPROT:017393; UNIPARC:UPI0000074F66; EMBL:AF026214; PIDN:AAB71313;
e: strain Bristol N2; clone F52H2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7-residue repeats
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            S07485; MUID:90098832;
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85.7%;
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85.7%;
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Zhou, L.; S
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 35; DB Pred. No. 9.8e 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                        <u>,</u>
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                                                                                                             from GB/EMBL/DDBJ
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        D.J.; Cummings, L.M.; Aravind, L.; Koonin, Sutton, G.G.; Clayton, R.; White, O.; Smith,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   predicted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
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72.3%;

Score

34;

BB

2

Length 140;

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A; McCesser.
A; Molecule type: DNA
A; Residues: 1-177 < DUB>
A; Cross-references: UNIPROT: P07249;
A; Cross-references: translated the (
                                                                                                                                                          C;Species: Saccharomyces cerevisiae
C;Date: 31-Dec-1991 #sequence revision 31-Dec-1991 #text_change 09-
C;Accession: S0582; S52891
R;Dubois, E.; Bercy, J.; Messenguy, F.
Mol. Gen. Genet. 207, 142-148, 1987
A;Title: Characterization of two genes, ARGRI and ARGRIII required A;Reference number: S05822; MUID:87257286; PMID:3298999
A;Accession: S05822; MUID:87257286; PMID:3298999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: UNIPARC:UPI000012F1D6; GB:M29712; NID:g205327; R;Vaughan, J.M.; Fischer, W.H.; Hoeger, C.; Rivier, J.; Vale, W. Endocrinology 125, 1660-1665, 1989
A;Title: Characterization of melanin-concentrating hormone from rat A;Reference number: A37406; MUID:89338286; PMID:2759038
A;Accession: A37406
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C,Date: 12-Apr-1991 Heequence revision
C,Accession: A36237, A37406
R;Thompson, R.C. Marron
                                                                     A;Note: the authors translated R;Odell, C.; Bowman, S.
                                                                                                                                                                                                                                                                                                           regulatory protein ARG80 - yeast (Saccharomyces cerevisiae)
N;Alternate names: protein YM9532.07; protein YMR042w; regu
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A; Residues: 1-165 < THO>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R;Thompson, R.C.; Watson, S.J. DNA Cell Biol. 9, 637-645, 1990
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A;Molecule
                                A; Reference number: S52885
                                                    submitted to the EMBL
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Matches 6
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;Residues: 147-165 <VAU>
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C; Function:
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A; Residues: 1-197 < LET>
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R;Le, T.
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                                                                        A;Cross-references: UNIPROT:Q40436; UNIPARC:UPI00000A96E4; GB:D26182; NID:g575607; PIDN C;Superfamily: wood tobacco RNA-binding protein; ribonucleoprotein repeat homology F;41-108/Domain: ribonucleoprotein repeat homology <RRM1>
                                                                                                                                                                                                                                                                                                  RNA-binding protein - wood tobacco
C;Species: Nicotiana sylvestris (wood tobacco)
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-
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A;Cross-references: SGD:S0004645; MIPS:YMR042w
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Mol. Gen. Genet. 244, 360-366, 1994
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probable serine rich protein - Arabidopsis thaliana
N;Alternate names: protein F14F18.220
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #text_C;Accession: T48564
R;Bevan, M.; Hilbert, H.; Braun, M.; Holzer, E.; Brandt,
submitted to the Protein
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A;Residues: 1-300 <KUB>
A;Cross-references: UNIPROT:O22090; UNIPARC:UPI00000ACA2C; EMBL:AB
A;Experimental source: strain Mitchell diploid
C;Superfamily: Arabidopsis thaliana hypothetical protein F12E4.290
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A;Reference number: 219885
A;Accession: T24387
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A;Title: Cys2/His2 zinc-finger protein family of petunia:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 zinc finger protein ZPT3-3 [imported] - garden petunia C;Species: Petunia x hybrida (garden petunia) C;Date: 24-Oct-2000 #sequence_revision 24-Oct-2000 #te: C;Accession: T52379
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A; Accession: T52379
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Status: preliminary; translated from
                                                                                                                                                                                                                                                                                                                      ;Introns; ;; papain ;; Superfamily: papain ;Keywords: cysteine proteinase; ;Keywords: 704/Active site: Cys, !
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;Status: preliminary; translated;Molecule type: DNA;Residues: 1-337 WIII>
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;Cross-references: UNIPROT:045734;
;Experimental source: clone T03E6
                                                                                                                                                                                                                                                                                                                                                                                   Introns: 86/3; 119/3; 269/1
                                                                                                                                                                                                                                                                                                                                                                                                                           Gene: CESP:T03E6.7
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Best Local
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                                                                                                                                                                                              DYKEDFDK 41
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 Sequence Database,
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75.0%;
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Pred. No. 1.7e
1; Mismatches
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                                                  20-Apr-2000 #text_change 09-Jul-2004
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 April 2000
                                                                                                                                                                                                                                                                                                  DB 2;
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1.5e+02;
1;
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                                                                                                                                                                                                                                                                                 .7e+02;
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                  A.;
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                  Duesterhoeft,
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 Genetics:
                                                                                                                                                                                                                                                                                                                                                                       Matches
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A;Cross-references: UNIPROT:D38199; UNIPARC:UPI000005318E; EMBL:Z35793; NID:g536038; A;Experimental source: strain S288C R;van Dyck, L.; Jonniaux, J.L.; de Melo Barreiros, T.; Kleine, K.; Goffeau, A. Yeast 10, 1633-1673, 1994 A;Title: Analysis of a 17.4 kb DNA segment of yeast chromosome II encompassing the ri human proliferation-associated p120 antigen. A;Reference number: S50299; MUID:95242843; PMID:7725803
                                                        A;Residues: 1-386 <BEV>
A;Cross-references: UNIPROT:O49617; UNIPARC:UPI00000AC134;
A;Experimental source: cultivar Columbia; BAC clone M4E13
                                                                                                                                                                            C;Accession: T05781
R;Bevan, M.; Purnelle, B.; Boutry, M.; Goffeau, A.; Hoheisel, submitted to the Protein Sequence Database, April 1998
                                                                                                                                                                                                                                           hypothetical protein M4E13.160 - Arabidopsis thaliana
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Datc: 23-Apr-1999 #sequence_revision 23-Apr-1999 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;ACCESSALO...
A;Molecule type: DNA
A;Residues: 1-381 <VAN>
A;Cross-references: UNIPARC:UPI00005318E;
A;Cross-references: Strain S288C
A; Map position: 4
A; Introns: 42/1;
                                                                                                                                            A;Reference number: 215451
A;Accession: T05781
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R;Goffeau, A.; Jonniaux, J.L.; Purnelle, B.; submitted to the Protein Sequence Database, FA;Reference number: S45745
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 hypothetical protein YBL032w - yeast (Saccharomyces cerevisiae) N;Alternate names: hypothetical protein YBL0418 C;Species: Saccharomyces cerevisiae
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A; Note: F14F18.220
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A;Residues: 1-362 <BEV>
A;Cross-references: UNIPROT:Q9LYH0; UNIPARC:UPI000009E7C1; EMBL:AL163812
A;Experimental source: cultivar Columbia; BAC clone F14F18
                                                                                                                    A; Molecule type: DNA
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A; Residues: 1-381 <GOF>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Date: 26-Aug-1994 #sequence_revision 09-Sep-1994 #text_change 09-Jul-2004
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                KDDDDK 302
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71.4%;
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No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Skala, J.;
August 1994
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                                                                                                                                                                                                           Mewes, H.W.;
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RESULT 56
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Experimental source: clone C;Genetics: A;Gene: CESP:Y48A6B.13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hypothetical protein Y48A6B.13 - Caenorhabditis elegans C;Species: Caenorhabditis elegans C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change C;Accession: T26987 R;Gardner, A. Species C,Accession: Data Library, June 1998 Submitted to the EMBL Data Library, June 1998
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T26987
probable calmodulin-binding protein [imported] -
C;Species: Arabidopsis thaliana (mouse-ear cress)
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A;Residues: 1-396 <WIL>
A;Cross_references: UNIPROT:Q9XXC7;
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A;Accession: T26987
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Best Local :
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tes 5; Conserv
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71.4%;
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   (mouse-ear cress)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2;
                                                                                                                                                                                 Score 34; DB 2;
Pred. No. 2.2e+02;
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Pred. No. 2e+02;
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Pred. No. 2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                  March
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                                                                                                                                                                                                                                                                                                                 UNIPARC: UPI000007D6C1; EMBL: Z70212; PIDN: CAA94163.1
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                 Arabidopsis thaliana
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                                                                                                                                                                                                Length 430;
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RESULT 57
S66713
                                                                                                                                           R;Delius, H.; Hofmann, B.
submitted to the EMBL Data Library, August 1993
A;Description: Primer-directed sequencing of human
                                                                                                                                                                                                                      B2 protein - human papillomavirus type 25
C;Species: human papillomavirus type 25
C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-467 <TTO>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Accession: A85363
R;anonymous, The European Union Arabidopsis Genome
Nature 402, 769-777, 1999
                                                                                                                                                                                                                                                                                                                                                                        뮍
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Gene: MIPS: YOL030w
A; Cross-references:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Molecule cype.
A;Residues: 1-484 <HAB>
A;Cross-references: UNIPROT: 008193;
A;Cross-references: strain $288C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Reference number: S66703
A; Accession: S66713
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R;Habbig, B.; Hattenhorst, U.; Hollenberg, C.P.; Ramezani submitted to the Protein Sequence Database, July 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hypothetical protein YOLO30w - yeast (Saccharomyces cerevisiae)
N,Alternate names: hypothetical protein O2145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Date: 16-Feb-2001 #sequence_revision 16-Feb-2001 #text_change 31-Dec-2004
                  A;Cross-references: UNIPROT:P36787; UNIPARC:UPI0000138340; EMBL:X74471; NID:g396948; C;Superfamily: papillomavirus E2 protein
                                                             A; Molecule type: DNA
A; Residues: 1-502 < DEL >
                                                                                                                     A; Reference number: S36469
                                                                                                                                                                                                        C; Accession: S36494
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Map position:
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                                                                                                     A; Accession: S36494
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Genetics:
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binding;
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  early protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         72.3%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 34; DB 2;
Pred. No. 2.5e+0;
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2; Mismatches
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  transcription regulation
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                                                                                                                                               papillomavirus
                                                                                                                                                                                                                           #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 484;
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Query Match

72.3%;

Score 34;

ВB

2;

Length 502

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RESULT 61
T13674
hypothetical protein EG0002.1 - fr
C;Species: Drosophila melanogaster
                                                                                                                                                                                                                                                                                                                                                                                                                        R;Kawagoe, Y.; Murai, N.
Plant Physiol. 117, 720, 1998
A;Title: PEP2, a second bHLH;
A;Reference number: Z17191
A;Accession: T10862
                                                                                                         밁
                                                                                                                                       δ
                                                                                                                                                                                                                                                                               A;Gene: PG2
C;Function:
                                                                                                                                                                                                                                                                                                                    A; Experimental source: C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              phaseolin G-box binding protein PG2 - )
C;Species: Phaseolus vulgaris (kidney k
C;Date: 16-Jul-1999 #sequence_revision
C;Accession: T10862
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Introns: 22/1; 58/1; 68/2
C; Superfamily. Schi---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R;Lyne, M.; Rajandream, M.A.; Barrell, B.G.; Rieger, M. submitted to the EMBL Data Library, November 1998 A;Reference number: Z21889 A;Accession: T40660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 59
T40660
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A;Residues: 1-609 <LYN>
A;Cross-references: UNIPROT:094326; UNIPARC:UPI000006C388; EMBL:AL034352; PIDN:CAA22180
A;Experimental source: strain 972h-; cosmid c725
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                                                                                                                                                                                           Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                      Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                  Status: preliminary; translated
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                                                                                                                                                                                                                                                Superfamily:
                                                                                                                                                                                                                                                                   Description:
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                                                                                                                                          YKDDDDK 8
                                                                                                         YKGDDDK 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           KDDDDK 364
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                                                                                                                                                                                                                                         transcription factor BHLH containing MYC-related DNA binding protein
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                                                                                                                                                                             Conservative
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                                                                                                                                                                                           72.3%;
85.7%;
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100.0%; Pred. No. 3.2e+02;
cive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein binding
                  fruit
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                                                                                                                                                                                           Score 34; DB 2;
Pred. No. 3.2e+0;
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); Mismatches 1
                                                                                                                                                                         Mismatches
                  fly
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                    (Drosophila
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                    melanogaster)
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                                                                                                R;Lin, X.; Kaul, S.; Rounsley, S.D.; Shea, T.P.; Benito, M.I.; Town, C.D.; Fujii, C.Y. M.; Koo, H.; Moffat, K.S.; Cronin, L.A.; Shen, M.; Vanaken, S.E.; Umayam, L.; Tallon, leuss, D.; Nierman, W.C.; White, O.; Eisen, J.A.; Salzberg, S.L.; Fraser, C.M.; Venter, Nature 402, 761-768, 1999
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-860 <ST
                                                 A;Reference number: A84420;
A;Accession: A84833
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                                                                                   A; Title: Sequence and analysis of chromosome
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<STO>

MUID:20083487;

me 2 of the plant PMID:10617197

Arabidopsis thaliana

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hypothetical protein At2940720 [imported] - Arabidopsis thaliana C;Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 02-Feb-2001 #sequence_revision 02-Feb-2001 #text_change 09-Jul-2004 C;Accession: A84833
                                                                                                                                                                                                                                                                                                                                                                                                   A;Cross-references: FlyBase:FBgn0025376
A;Introns: 27/3; 55/1; 131/2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hypothetical protein EG0002.1 - fruit fly (Drosophila melanogaster) C;Species: Drosophila melanogaster C;Species: 13-Aug-1999 #sequence_revision 13-Aug-1999 #text_change 09-
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R;Bolshakov, V.; Borkova, D.; Minana, B.; Kafatos, F. submitted to the EMBL Data Library, September 1998 A;Description: Sequencing the distal X chromosome of Drosophila melanogaster A;Reference number: Z17698
                                                                                                                                                                                                                                                                                                                                                                                  A;Note: EG:EG0002.1
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A; Residues: 1-743 <BOL>
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R; Bolshakov, V.; Bo
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A;Note: EG:EG0002.1
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A; Residues: 1-734 <BOL>
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Matches
                                                                                                                                                                                                                                                                                                                                      ;Superfamily: peptide transport;Keywords: alternative splicing
                                                                                                                                                                                                                                                                                               Query Match
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EMBL Data Library, September 1998
guencing the distal X chromosome (
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Pred. No. 3.9e+02;
2; Mismatches 0;
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Pred. No. 3.9e+02
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C;Accession: B96508
R;Theologis, A; Ecker, J.R.; Palm, C.J.; Federspiel, N.A.; Kaul, S.; White, O.; Alonso Chin, C.W.; Chung, M.K.; Conn, L.; Conway, A.B.; Conway, A.R.; Creasy, T.H.; Dewar, K. Nature 408, 816-820, 2000
A;Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khaykin, E.; Kim, C.C.A.; Li, J.H.; Li, Y.; Lin, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Maiti, R.; Marziali Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.
A;Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sun, H.; Tallon, ker, M.; Wu, D.; Yu, G.; Fraser, C.M.; Venter, J.C.; Davis, R.W.
A;Title: Sequence and analysis of chromosome 1 of the plant Arabidopsis.
A;Reference number: A86141; MUID:21016719; PMID:11130712
                                                                                                                                                                                                                                                         C;Accession: A41996
R;Capobianco, A.J.; Chang, D.; Mosialos, G.; Gilmore, T.D.
J. Virol. 66, 3758-3767, 1992
A;Title: p105, the NF-kappa-B p50 precursor protein, is on A;Reference number: A41996; MUID:92260650; PMID:1533881
A;Accession: A41996
                                                                                                 C;Superfamily: human transcription of C;Keywords: DNA binding; nucleus; pt C;Keywords: DNA binding; nucleus; pt F;47-371/Domain: rel homology cRE;
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C;Genetics:
A;Gene: At2g40720
                                                                       F;366-370/Region: nuclear location signal F;342/Binding site: phosphate (Ser) (cova
                                                                                                                                                                                                                                                                                                                                                                                               NF-kappa-B p50 subunit precursor p105 - chicken
C;Species: Gallus gallus (chicken)
C;Date: 06-Nov-1992 #sequence_revision 06-Nov-1992 #text_change 09-Jul-2004
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C; Superfamily: replication licensing factor MCM2; MCM homology
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C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 09-Jul-2004
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                                                                                                                                                                                               ;Status: preliminary
;Molecule type: mRNA
;Residues: 1-984 <CAP>
                                                                                                                            Cross-references: UNIPROT:Q04861; UNIPARC:UPI000012DC52; GB:M86930; NID:g212525; PIDN;Superfamily: human transcription factor NF-kappa-B2; ankyrin repeat homology; rel hom;Keywords: DNA binding; nucleus; phosphoprotein; transcription regulation
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                                                                   site: phosphate (Ser)
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Pred. No. 4.6e
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Score 34;
Pred. No.
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Pred. No. 5e+02;
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. 4.6e+02;
  5.3e+02;
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                       Length 984;
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R;Theologis, A.; Ecker, J.R.; Palm, C.J.; Federspiel, N.A.; Kaul, S.; White, O. Chin, C.W.; Chung, M.K.; Conn, L.; Conway, A.B.; Conway, A.R.; Creasy, T.H.; I ansen, N.F.; Hughes, B.; Huizar, L.
Nature 408, 816-820, 2000
A;Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khaykin, B. A;Authors: Hunter, J.L.; Lin, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Maiti, R.; C.A.; Li, J.H.; Li, Y.; Lin, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Maiti, R.; Rizzo, M.; Rooney, T.; Rowley, D.; Sakkano, H.
A;Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sun, H.; ker, M.; Wu, D.; Yu, G.; Fraser, C.M.; Venter, J.C.; Davis, R.W.
A;Title: Sequence and analysis of Chromosome 1 of the plant Arabidopsis.
A;Reference number: A86141; MUID:21016719; PMID:11130712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 phospholipase C - yeast (Candida albicans)
C;Species: Candida albicans
C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 09-Jul-2004
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F;566-726/Domain: 1-phosphatidylinositol-4,5-bisphosphate phosphodiesterase domain X ho
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A; Cross-references: UNIP
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A; Reference number: Z18844; MUID:98129081; PMID:9467900
                                                                                                                                                                                                                                                                                                                          A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                A; Reference number: A86141; A; Accession: C96761
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hypothetical protein T9L24.35 [imported] - Arabidopsis thaliana
C;Species: Arabidopsis thaliana (mouse-ear cress)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Accession: T18257
                                                                                                                                                                                                                                                      ;Residues: 1-1157 <STO>
;Cross-references: UNIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Accession: C96761
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                          Gene: T9L24.35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           628 KDDDDK 633
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       757
421 EYEDDDD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             D.E.; Mccreary, C.E.; Coleman, D.C.
gy 144, 55-72, 1998
                                                                                     l Similarity
5; Conserv
                                       1 DYKDDDD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3 KDDDDK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DVKDDDD 763
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                       Conservative
                                                                                                                                                                                                                                                                                                    DNA
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                                                                                                                                                                                                                                                      UNIPROT:Q9FX39; UNIPARC:UPI000009E7EF; GB:AE005173; NID:g11120792;
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427
                                                                                                                72.3%;
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                                                                                          Score 34; DB
Pred. No. 6.2e
2; Mismatches
                                                                                          <u>۲</u>.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C.J.; Federspiel, N.A.; Kaul, S.; White, O.; Alonso Conway, A.B.; Conway, A.R.; Creasy, T.H.; Dewar, K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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, 6.2e+02;
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beta prime

chain [imported] -

Nostoc sp.

(strain PCC

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A; Molecule type: DNA
A; Residues: 1-937, 'A', 939-1429, 'C', 1431-1953 < FAR>
A; Residues: 1-937, 'A', 939-1429, 'C', 1431-1953 < FAR>
A; Cross-references: UNIPARC: UPI0000168E7F; EMBL: L31766; NID: 9472524; PIDN: AAA34455.1;
R; Sen-Gupta, M.; Lyck, R.; Fleig, U.; Niedenthal, R.N.; Hegemann, J.M.
R; Sen-Gupta, M.; Lyck, R.; Fleig, U.; Niedenthal, R.N.; Hegemann, J.M.
R; Sen-Gupta, M.; Lyck, R.; Fleig, U.; Niedenthal, R.N.; Hegemann, J.M.
R; Sen-Gupta, M.; Lyck, R.; Fleig, U.; Niedenthal, R.N.; Hegemann, J.M.
R; Sen-Gupta, M.; Lyck, R.; Fleig, U.; Niedenthal, R.N.; Hegemann, J.M.
R; Sen-Gupta, M.; Lyck, R.; Fleig, U.; Niedenthal, R.N.; Hegemann, J.M.
R; Sen-Gupta, M.; Lyck, R.; Fleig, U.; Niedenthal, R.N.; Hegemann, J.M.
R; Sen-Gupta, M.; Lyck, R.; Fleig, U.; Niedenthal, R.N.; Hegemann, J.M.
R; Sen-Gupta, M.; Lyck, R.; Fleig, U.; Niedenthal, R.N.; Hegemann, J.M.
R; Sen-Gupta, M.; Lyck, R.; Fleig, U.; Niedenthal, R.N.; Hegemann, J.M.
R; Sen-Gupta, M.; Lyck, R.; Fleig, U.; Niedenthal, R.N.; Hegemann, J.M.
R; Sen-Gupta, M.; Lyck, R.; Fleig, U.; Niedenthal, R.N.; Hegemann, J.M.
R; Sen-Gupta, M.; Lyck, R.; Fleig, U.; Niedenthal, R.N.; Hegemann, J.M.
R; Sen-Gupta, M.; Lyck, R.; Fleig, U.; Niedenthal, R.N.; Hegemann, J.M.
R; Sen-Gupta, M.; Lyck, R.; Fleig, U.; Niedenthal, R.N.; Hegemann, J.M.
R; Sen-Gupta, M.; Lyck, R.; Fleig, U.; Niedenthal, R.N.; Hegemann, J.M.
R; Sen-Gupta, R. R; S
                                      Yeast 12, 505-514, 1996
A;Title: The sequence of a 24 152 bp segment from the left A;Reference number: 865111; MUID:96310631; PMID:8740425 A;Accession: S65111
                                                                                                                                               A;Reference number: S60909
A;Accession: S60909
A;Molecule type: DNA
A;Residues: 1-1553 <SE2>
A;Cross-references: UNIPARC:UPI0000168B2A; EMBL:X92494;
A;Cross-references: UNIPARC:UPI0000168B2A; EMBL:X92494;
A;Cross-Cupta, M.; Lyck, R.; Fleig, U.; Niedenthal, R.K.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Species: Nostoc sp. PCC 7120
A;Note: Nostoc sp. strain PCC 7120 is a synonym of Anabaena sp. C;Date: 14-Dec-2001 #sequence_revision 14-Dec-2001 #text_change C;Accession: AF2005
R;Kaneko, T.; Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S. Nakazaki, N.; Shimpo, S.; Sugimoto, M.; Takazawa, M.; Yamada,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Residues: 1-1553 <SEN>
A;Rcosb-references: UNIFART: P41832; UNIPARC: UPI0000168B2A; EMBL: Z71547; MIPS: YNL;Z71c A;Cxosb-references: Strain S288C
A;Experimental source: strain S288C
R;Messenguy, F.; Dubois, E.; Vierendeels, F.; Scherens, B.; Pierard, A.; Glansdorff, submitted to the Protein Sequence Database, April 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C;Date: 27-Apr-1996 #sequence revision 03-May-1996 #text_change 09-Jul-2004 C;Accession: S63244; S63245; S48523; S60909; S65111 R;Sen-Gupta, M.; Lyck, R.; Niedenthal, R.K.; Fleig, U.N.; Hegemann, J.H. submitted to the Protein Sequence Database, April 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Fares, H.F.; Pringle, J.R.
submitted to the EMBL Data Library,
A;Description: Synthetic Lethals of
A;Reference number: S48524
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Molecule type: DNA
A; Residues: 987-1953 <MES>
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A; Accession: S63245
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA Res. 8, 205-213, 2001
A;Title: Complete Genomic Sequence of the Filamentous Nitrogen-fixing Cyanobacterium A;Reference number: AB1807; MUID:21595285; PMID:11759840
A;Accession: AF2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Accession: S48523
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Experimental source: strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: UNIPARC:UPI000017B262; EMBL:Z71547; MIPS:YNL271c
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C;Superfamily: chloroplast DNA-directed
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A;Molecule type: DNA
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Best Local S
Matches 5
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5; Conserv
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71.4%;
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Pred. No.
      shown;
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CDC12.
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7.3e+02;
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                                                                                                                                                         NID:g1045236;
Hegemann, J.H.
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                                                                                                                                                                                      PIDN: CAA63225.1;
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A; Accession: S76216
A; Status: nucleic ac
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A;Variety: PCC 6803
C;Date: 25-Apr-1997 #seqn
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                                                                                                                                                                                                                      A;Start
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Title: Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Note:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Map position:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Cross-references:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Residues: 1-4550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Status:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Reference number: A; Accession: T18440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C;Accession: T18440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                T18440
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                                                                                                                            Query Match
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Best Local
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                                                                                                                         Local
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5; Conser
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62.
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A; Nolecule type: DNA
A; Residues: 1-125 <KAN>
A; Residues: 1-125 <KAN>
Norman-references: UNIPROT: P74379; UNIPARC: UPI00000C0E6F; EMBL: D90914;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C;Accession: S76216
R;Kaneko, T.; Sato, S.; Kotani, H.; Tanaka, A.; Asar
R;Kaneko, T.; Sato, S.; Shimpo, S.; Takeuchi, C.; Wada,
DNA Res. 3, 109-136, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hypothetical protein C0425w - malaria parasite (Plasmodium falciparum) C;Species: Plasmodium falciparum C;Species: Plasmodium falciparum C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 09-Jul
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Gene: SGD:BNI1; SHE5; SYL39
A;Cross-references: SGD:S0005215; MIPS:YNL271c
A;Map position: 14L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Modecule type: DNA
A;Modecule type: DNA
A;Residues: 1-1553 <SEW>
A;Residues: 1-1553 <SEW

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       preliminary; translated
                                                                                                                                                                                                                                                                                                                                                                             acid sequence not shown; translation not shown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           analysis of the genome of the unicellular cyanobacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                            S74322; MUID:97061201;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  #sequence_revision 25-Apr-1997 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                UNIPROT:077336;
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75.0%;
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Pred. No. 2.6e
1; Mismatches
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                                   Score 33;
Pred. No.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           from
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                                           B8;
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                                                                       2;
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<u>;</u>
                                                                   Length 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           T.; Watanabe,
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                                                                                                                                                                                                                          June 1996
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da, M.; Yasur
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C; Date: (C; Accessi
R; Smith,
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                                                                                                                                                                                                                                                                                                            A; Molecule type: DNA
A; Residues: 1-147 < DOO>
                                                                                                                                                                                                                                                                                                                                           A;Title: Sequence divergence
A;Reference number: A46315; N
A;Accession: B46315
                                                                                                                                                                                                                                                                                                                                                                                                      R; Doorbar,
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C;Function:
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C;Species: Dermatophagoides farinae
C;Date: 28-Aug-1985 #sequence revision 07-Oct-1994 #text_change 31-Dec-2004
C;Accession: JX0313; PC2125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          밁
              hypothetical protein MTH1578 - Methanobacterium thermoautotrophicum (strain c;Species: Methanobacterium thermoautotrophicum C;Date: 05-Dec-1997 #sequence_revision 05-Dec-1997 #text_change 09-Jul-2004 C;Accession: E69077
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                                                                                                                                                                                                                                                                                                                                                                                          Virology 172, 51-62,
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                                                                                             RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                   4 protein - human papillomavirus type ;Species: human papillomavirus type 4 ;Date: 31-Dec-1993 #sequence_revision ;Accession: B46315
                                                                                                                                                                                                                    Best
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Reference number: JX0313; MUID:94334283; PMID:8056755
                                                                                                                                                                                                                                                                   Keywords:
                                                                                                                                                                                                                                                                                 Superfamily: papillomavirus type 4
                                                                                                                                                                                                                                                                                              Cross-references: UNIPARC:UPI00000F4EF0; GB:M28744;
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                                                                                                                                                                                                                    Local
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 D.R.;
                                                                                                                                                                                                                                                                                                                                                                                                  J.;
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                                                                                                                                          DYDEDDEK
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   Doucette-Stamm,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      435-440,
                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                           1989
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                                                                                                                                          85
                                                                                                                                                                                                                                                                                                                                                                                                      I.; Gallimore,
                                                                                                                                                                                                                    70.2%;
62.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         70.2%;
                                                                                                                                                                                                                                                                                                                                                           yet conserved
MUID:89370332;
 L.A.; Deloughery,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 33;
Pred. No.
                                                                                                                                                                                                                  Score 33; DB 1;
Pred. No. 1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            UNIPARC: UPI000012EAEB; DDBJ:D17676; NID:g642918; this amino acid sequence is inconsistent with the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                              E4 protein
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                                                                                                                                                                                                                                                                                                                                                                                                          P.H
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                                                                                                                                                                                                                                                                                                                                                             physical characteristics PMID:2549722
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1e+02;
 C.; Lee,
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                                                                                                                                                                                                                                  Length 147;
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                                                                                                                                                                                                                                                                                               NID:g333136;
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   H.;
 Dubois,
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S59766
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   Matches
                 Query Match
Best Local :
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; Qiu, D.; Spadafora, R.; Vicaire, R.; Wang, Y.; Wierzbowski, J.; Gibson, R.; Jiwani, ki, S.; Church, G.M.; Daniels, C.J.; Mao, J.; Rice, P.; Noelling, J.; Reeve, J.N. J. Bacteriol. 179, 7135-7155, 1997
A;Title: Complete genome sequence of Methanobacterium thermoautotrophicum Delta H: fur A;Reference number: A69000; MUID:98037514; PMID:9371463
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Map position:
C; Superfamily:
                                                                                                                                                                                                  C;Accession: A61442
R;Amons, R.; Schipper, A.; van
J. Protein Chem. 11, 404, 1992
                                                                                                                                                                                                                                                          translation elongation factor eEF-1 beta chain - pig (fragments)
C;Species: Sus scrofa domestica (domestic pig)
C;Date: 07-Oct-1994 #sequence_revision 07-Oct-1994 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R;Nelson, ... submitted to the EMBL Data Library, July 175.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hypothetical protein YPR101w - yeast (Saccharomyces N;Alternate names: hypothetical protein P8283.11
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A;Gene: MTH1578
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A;Experimental source: strain Delta
                                                                   A; Molecule type: protein
A; Residues: 1-188 < AMO>
A; Cross-references: UNIPROT: Q7M397; UNIPARC: UPI00001782A2
                                                                                                                                                                A; Reference number: A61442
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: DNA
A; Residues: 1-175 <NEL>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Molecule type: DNA
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                                                                                                                             A;Status: preliminary
                                                                                                                                                A; Accession: A61442
                                                                                                                                                                                     A; Title: The primary structure of elongation factor lbeta from
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;Species: Saccharomyces cerevisiae;Date: 13-Jan-1996 #sequence_revision 01-Mar-1996
                                                       Superfamily: translation elongation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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 Score 33;
Pred. No.
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Pred. No. 1
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                     Length 188;
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                                                         chain;
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Similarity 6; Conserv

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N;Alternate names: protein YAL014c
C;Species: Saccharomyces cerevisiae
C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 09-Jul-2004
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                                                                                                                                                                                                                                                                                                      A; Description: Sequencing of A; Reference number: $36711
                                                                                                                                                                                                                                                                                                                                               C;Accession: S36720
R;Ouellette, F.; Clark, M.W.; Keng, submitted to the EMBL Data Library,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Residues: 1-190 < YOS>
A;Cross-references: UNIPROT:P78859; UNIPARC:UPI000013A6A4; |
A;Cross-references: UNIPROT:P78859; UNIPARC:UPI000013A6A4; |
A;Experimental source: strain PR745
R;Wood, V.; Rajandream, M.A.; Barrell, B.G.; Rieger, M.
submitted to the EMBL Data Library, March 1999
A;Reference number: Z22000
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T43013
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       conserved hypothetical protein SPCC645.03c - fission yeast (Schizosaccharomyces C;Species: Schizosaccharomyces pombe
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A;Accession: T43013
                                                                                                                                                                                                                          Cross-references: UNIPROT: P31377; UNIPARC: UPI000013A006;
                                                                                                                                                                                                                                                   Residues: 1-205 <0
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;Residues: 1-190 <WOO>
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Best Local
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71.4%;
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83.3%;
                                                                                                                                                                                                                                                                                                                          ; Keng, T.; Storms, R.K.; Zhong, W.; Zeng, B.; Fortin, N.; ibrary, January 1993
Chromosome I from Saccharomyces cerevisiae: analysis of a
                                                                     Score 33; DB
Pred. No. 1.5e
2; Mismatches
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Pred. No. 1.4e+02;
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1.5e+02;
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translation elongation factor eEF-1 beta chain - rate; C;Species: CryCtclagus cuniculus (domestic rabbit) C;Date: 14-Feb-1997 #sequence revision 13-Mar-1997 C;Accession: $62693; $62688; $37087 R;Chen, C.J.; Traugh, J.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R;Sanders, J.; Maassen, J.A.; Amons, R.; Moeller, W.
Nucleic Acids Res. 19, 4551, 1991
A;Title: Nucleotide sequence of human elongation factor-1-beta
A;Reference number: S25432; MUID:91360360; PMID:1886777
A;Accession: S25432
submitted to the EMBL Data Library, August 1993 A;Reference number: S37087
                                                                                                                                                                                                                                                                                                        Biochim. Biophys. Acta 1264, 303-311, 1995
A;Title: Expression of recombinant elongation factor 1
A;Reference number: $62688; MUID:96138552; PMID:8547318
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C;Superfamily: translation elongation factor eEF-1 beta chain; t
C;Keywords: phosphoprotein; protein biosynthesis
F;2-225/Product: translation elongation factor eEF-1 beta chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Experimental source: ovarian granulosa cells R;Bauw, G.; Rasmussen, H.H.; Van Den Bulcke, M. Electrophoresis 11, 528-536, 1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R;Von der Kammer, H.; Klaudiny, J.; Zimmer, M.; Scheit, K.H. Biochem. Biophys. Res. Commun. 177, 312-317, 1991
A;Title: Human elongation factor 1beta: cDNA and derived amino A;Reference number: JQ1038; MUID:91254293; PMID:1710449
A;Accession: JQ1038
                                                                                   A; Cross-references:
                                                                                                              A;Molecule type: protein
A;Residues: 80-91 <CHF>
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A;Residues: 2-6,'X','X',56-60 <BAU>
A;Cross-references: UNIPARC:UPI0000174808; UNIPARC:UPI0000174809
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                                                                                                                                                                          A;Accession: S62688
                                                                                                                                                                                                     A;Cross-references:
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A; Residues: 1-225 < VON>
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R;Von der Kammer, H.; Klaudiny, J.; Zimmer, M.; Scheit, K.H.
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A; Residues: 1-225 <SAN>
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                                                                                   UNIPARC: UPI00001782A7
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85.7%;
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Pred. No. 1.6e+02;
0; Mismatches 1; Indels
                                                                                                                                                                                                  UNIPARC: UPI00001782A6; EMBL: X74728
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PID: в. ;

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A; Molecule type: mRNA
A; Residues: 1-77, 'T', 79-225 < CHW>
A; Cross-references: UNIPARC: UPI000
C; Genetics:
A; Gene: EFIB
C; Function:
A; Pathway: protein biosynthesis
C; Superfamily: translation elongat
C; Keywords: GTP exchange; phosphot
F; 2-225/Product: translation elong
F; 97-105/Region: beta, beta' and of
F; 106/Binding site: phosphate (Se)
                                                                                                                                                                                                                                                                      R;Lin, X.; Kaul, S.; Rounsley, S.D.; Shea, T.P.; Benito, M.I.; Town, C.D.; Fujii, C.Y. M.; Koo, H.; Moffat, K.S.; Cronin, L.A.; Shen, M.; VanAken, S.E.; Umayam, L.; Tallon, euss, D.; Nierman, W.C.; White, O.; Eisen, J.A.; Salzberg, S.L.; Fraser, C.M.; Venter,
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A; Residues: 1-235 < KUR>
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A;Reference number: A99512; MUID:21267165; PMID:11353084
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Chambaud, I.; Heilig, R.; Ferris, S.; Barbe, Nucleic Acids Res. 29, 2145-2153, 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hypothetical protein MYPU_1110 [imported] - Mycoplasma pulmonis C;Species: Mycoplasma pulmonis
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       A;Gene: At2g4668
A;Map position:
C;Keywords: DNA
                                                                      A; Cross-references: C; Genetics:
                                                                                                                                      A; Molecule type: DNA
                                                                                                                                                           A; Status: preliminary
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                                                                                                                                                                                   Reference number: A84420; Accession: H84905
                                                                                                                                                                                                             ;Title: Sequence and analysis of chromosome 2 of the plant Arabidopsis thaliana.;Reference number: A84420; MUID:20083487; PMID:10617197
                                                                                                                                                                                                                                                                                                                                             Date: 02-Feb-2001 #sequence_revision 02-Feb-2001 #text_change 05-Oct-2004; Accession: H84905
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Date: 24-May-2001 #sequence_revision 24-May-2001 #text_change
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Cross-references: UNIPARC:UPI000016C566; EMBL:X74728; NID:g398394; PIDN:CAA52741.1;
                                                                                                                      ;Residues: 1-258 <STO>
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Best Local
                                                 Gene: At2g46680
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binding; homeobox; nucleus;
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                                                                                            UNIPROT: P46897;
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85.7%;
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71.4%;
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Pred. No. 1.7e
2; Mismatches
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                                                                                            UNIPARC: UPI0000126233;
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    transcription
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    regulation
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A;Cross-references: UNIPROT:P21828; UNIPARC:UPI000012A56D; R;Kikuchi, Y.; Mori, K.; Suzuki, S.; Yamaguchi, K.; Mizuno Gene 110, 151-158, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Note: the authors translated the codon CGA for residue 46 as Ala R;Mori, K.; Kikuchi, Y.; Honma-Waga, M.; Waga, S.; Mizuno, S. submitted to the EMBL Data Library, October 1993
A;Description: Production of a chimeric fibroin light-chain polypeptide as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R;Yamaguchi, K.; Kikuchi, Y.; Takagi, T.; Kikuchi, A.; Oyan
J. Mol. Biol. 210, 127-139, 1989
A;Tille: Primary structure of the silk fibroin light chain
A;Reference number: A33470; MUID:90064541; PMID:2585514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C;Date: 27-Feb-1990 #sequence revision 27-Feb-1990 #text_change 09-Jul-2004 C;Accession: A33470; JH0580; S41445; S41446; S10622 R;Yamaguchi, K.; Kikuchi, Y.; Takagi, T.; Kikuchi, A.; Oyama, F.; Shimura,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Molecule type: DNA
A;Acesidues: 1-45,'k',47-262 <KIK>
A;Residues: 1-145,'k',47-262 <KIK>
A;Cross-references: UNIPARC:UPIO00016B892; GB:M76430; NID:g289362; PIDN:AAA27840.1;
A;Experimental source: strain J-139
A;Experimental source: strain J-139
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Molecule type: mRNA
A; Residues: 1-262 < YA
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                                                                                                                                                                                                                                                                                                       A; Molecule type:
A; Residues: 1-12
                                                                                                                                                                                                                                                                                                                                                   A; Reference number: S10622; A; Accession: S10622
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A; Residues: 1-105, 'SMO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Accession: A33470
                                                                                           F;17-26/Product:
                                                                                                                                                                                                                                  A;Note: the complete translation is C;Comment: Silk fibroin is secreted
                                                                                                                                                                                                                                                                                 A;Cross-references: UNIPARC:UPI0000178C89; GB:X53319; NID:g10893
                                                                                                                                                                                                                                                                                                                                                                                 A; Title: Homeodomain binding sites in the ! A; Reference number: S10622; MUID: 90278949;
                                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: UNIPARC:UPI000007794F;
A;Experimental source: Nd-sD mutant
R;Hui, C; Suzuki, Y; Kikuchi, Y; Mizuno,
J. Mol. Biol. 213, 395-398, 1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Molecule type:
A;Residues: 1-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Experimental source: Nd-s mutant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Cross-references: UNIPARC:UPI0000077132;
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Matches
                                                                                                                                    Keywords: extracellular protein
                                                                                           .6/Domain: signal sequence #status predicted <SIG>.26/Product: fibroin light chain #status predicted
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6
                 Similarity
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                                                                                                                                                                                                                                                                                                            <HUI>
                                                                                                                                                                                                                                                                                                                                  DNA
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71.4%;
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75.0%;
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Pred. No. 1.9e
2; Mismatches
  0,
                        Score 33; DB 2;
Pred. No. 1.9e+02;
                                                                                                                                                                                                                                     not shown from the posterior
Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                 5' flanking region
PMID:1972197
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                                              DB 2;
                                                                                           predicted
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                                            Length 262;
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  Indels
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  Gaps
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R;Theologis, A.; Ecker, J.R.; Palm, C.J.; Federspiel, N.A.; Kaul, S.; White, Chin, C.W.; Chung, M.K.; Conn, L.; Conway, A.B.; Conway, A.R.; Creasy, T.H.; ansen, N.F.; Hughes, B.; Huizar, L.

Nature 408, 816-820, 2000
A;Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khaykin, E.C.A.; Li, J.H.; Lin, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Maiti, R.; C.A.; Li, Y.; Lin, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Maiti, R.; Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.

A;Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sun, H.

A;Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sun, H.

ker, M.; Wu, D.; Yu, G.; Fraser, C.M.; Venter, J.C.; Davis, R.W.

A;Reference number: A86141; MUID:21016719; PMID:11130712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Cross-references: UNIPARC:UPI000017A2CA;
C;Keywords: DNA binding; homeobox; leucine
F;39-95/Domain bearing
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RESULT
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A; Residues: 37-45,'N',47-133 <SOF>
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Plant Mol. Biol. 26, 145-154, 1994
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A; Residues: 1-267 < SOE>
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C;Species: Arabidopsis thaliana (mouse-ear cre
C;Date: 13-Jan-1995 #sequence_revision 13-Jan-
                                                                                                                                                                                                                                                                                                                          Status: preliminary
                                                                                                                                                                                                                                                                                                                                              Reference number:
                                                                                                                                                   Query Match
Best Local
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Best Local
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                                                                                                                                                                                                            position: 1
  86
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71.4%;
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71.4%;
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Library, June 1992
                                                                                                                           Score 33; DB:
Pred. No. 2e+0:
2; Mismatches
                                                                                                                               2
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zipper; nucleus; transcription
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2e+02;
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Maiti, R.;
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war, K.;
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C;Accession:
R;Beyer, S.;
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Plant J. 16, 263-276, 1998
A;Title: Towards functional
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T51667
                                    R;Beyer,
                                                                                                                                                                                                  S49469
                                                                                                                                                                                                                                   RESULT 88
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C;Superfamily: Myb-related transcription activator;
C;Keywords: transcription factor
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                submitted to
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A;Experimental source: cultivar Columbia
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A; Accession: T51667
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submitted to the EMBL Data Library, April 1998
a.Deacription: Genomic sequence for Arabidopsis thaliana
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A;Cross-references: UNIPROT:Q8GYP5; UNIPARC:UPI000000C547; EMBL:AC003981; NID:g3063438
A;Experimental source: cultivar Columbia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;Superfamily: Myb-related transcription activator; myb;9-61/Domain: myb DNA-binding repeat homology <MYB1>;62-112/Domain: myb DNA-binding repeat homology <MYB>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gene: MYB60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Status: preliminary; translated from GB/EMBL/DDBJ
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Best Local
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                                                                                                                                                                                                                                                                                                                               236 DHEDDDD 242
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5; Conserv
                the EMBL
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.M.; Piepersberg, W.
Data Library, October 1994
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71.4%;
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Pred. No.
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Pred. No. 2.1e+02;
2; Mismatches 0;
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                                                                                                                                                             Streptomyces glaucescens
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                                                                                                      09-Jul-2004
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R;Palmer, S.
submitted to the EMBL Data Library
A;Reference number: Z19213
A;Accession: T20033
A;Status: preliminary; translated
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R;Tettelin, H.; Nelson, K.E.; Paulsen, I.T.; Eisen, J.A.; Read, T.D.; Peterson, S.; Heicon, J.D.; Umayam, L.A.; White, O.; Salzberg, S.L.; Lewis, M.R.; Radune, D.; Holtzapple, nson, T.; Hickey, E.K.; Holt, I.E.

Science 293, 498-506, 2001

A;Authors: Loftus, B.J.; Yang, F.; Smith, H.O.; Venter, J.C.; Dougherty, B.A.; Morrison, A;Title: Complete Genome Sequence of a virulent isolate of Streptococcus pneumoniae.

A;Reference number: A95000; MUID:21357209; PMID:11463916

A;Accession: G95244
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G95244
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A;Accession: S49469
A;Atacus: preliminary
A;Molecule type: DNA
A;Residues: 1-299 <AEY>
A;Cross-references: UNIPROT:Q54266; UNIPARC:UPI00000BA7DC; EMBL:X82177; NID:g902359; PII
C;Superfamily: glucose-1-phosphate cytidylyltransferase
C;Keywords: nucleotidyltransferase
                                                                                                                                                                                                                                                                                                                        C;Species: Caenorhal
C;Date: 15-Oct-1999
C;Accession: T20033
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hypothetical protein SP2093 [imported] - Streptococcus pneumoniae (strain TIGR4) C;Species: Streptococcus pneumoniae C;Decies: Streptococcus pneumoniae C;Date: 03-Aug-2001 #sequence_revision 03-Aug-2001 #text_change 09-Jul-2004 C;Accession: G95244
                                                                                                                                                                                         A; Molecule type: DNA
A; Residues: 1-333 <WIL>
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A;Experimental source: strain TIGR4
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                                                                                                                                                      Cross-references: UNIPROT:Q18694; Experimental source: clone C47G2
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Best Local :
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Best Local (
                                                                                                                   Gene: CESP:C47G2.2
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                                                                            position: 2
rons: 113/2; 222/2
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ص
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75.0%;
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85.7%;
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Pred. No. 2.4e
0; Mismatches
     Score 33; DB
Pred. No. 2.5e
0; Mismatches
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Pred. No. 2.2e+02;
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                                                                                                                                                                       UNIPARC: UPI0000080191; EMBL: Z49125; PIDN: CAA88935.1
                                                                                                                                                                                                                                                                                     April
                                                                                                                                                                                                                                  GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                           15-Oct-1999 #text_change 09-Jul-2004
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2;
..5e+02;
1;
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                                       Length 333;
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Science 294, 849-852, 2001
Science 294, 849-852, 2001
A;Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; Madueno, A;Authors: Kreft, T.; Simoes, N.; Tierrez, A.; Vazquez-Boland, J. ok, C.; Schlueter, T.; Simoes, N.; Tierrez, A.; Vazquez-Boland, J. A;Title: Comparative genomics of Listeria species.
A;Title: Comparative genomics of Listeria species.
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cell wall binding proteins homolog lmo2504 [imported] - Listeria monocytogenes C; Species: Listeria monocytogenes C; Species: Listeria monocytogenes C; Date: 27-Nov-2001 #sequence_revision 27-Nov-2001 #text_change 09-Jul-2004 C; Accession: AH1387
C; Accession: AH1387
R; Glaser, P.; Frangeul, L.; Buchrieser, C.; Amend, A.; Baquero, F.; Berche, P.; R; Glaser, P.; Frangeul, G.; Duchaud, E.; Durand, L.; Dussurget, O.; Entian, K.D.; Dominguez-Bernal, G.; Duchaud, E.; Durand, L.; Dussurget, O.; Entian, K.D.;
                                                                                                                     RESULT 93
T40337
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Weisshaar, B.; Armstrong, G.A.; Block, A.; da Costa e Silva, O.; Hahlbrock, EMBO J. 10, 1777-1786, 1991
A;Title: Light-inducible and constitutively expressed DNA-binding proteins re A;Reference number: S16320; MUID:91266906; PMID:2050115
                      probable saccharopine dehydrogenase (NADP, L-glutamate-forming) (EC 1.5.1.10) C;Species: Schizosaccharomyces pombe C;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 09-Jul-2004 C;Accession: T40337
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A;Molecule type: mRNA
A;Residues: 1-393 <WEI>
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C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 31-Dec-2004
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C;Accession: T40337
R;Lyne, M.; Rajandream, M.A.; Barrell, B.G.; Beck,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: DNA
A; Residues: 1-436 <GLA>
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Similarity 62.5%;
5; Conservation
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75.0%;
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Pred. No. 3.3e+02;
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     R.; Pohl,
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Voss, H.; Wehland
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Fsihi, H
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submitted to the

EMBL Data Library,

March 1998

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$26606

myb-related protein 3 - garden petunia
C:Species: Petunia x hybrida (garden petunia)
C;Date: 25-Feb-1994 #sequence_revision 03-Nov
C;Accession: $26606
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Keywords: heme; iron; metalloprotein; oxidoreductase F;59,99,132/Active site: His, Thr, Asn #status predicted F;342/Binding site: heme iron (Tyr) (axial ligand) #stat
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A;Residues: 1-504 <TET>
A;Residues: 1-504 <TET>
A;Cross-references: UNIPROT:Q9JRF5; UNIPARC:UPI00000C4DD7; GB:AE002379;
A;Cross-references: UNIPROT:Q9JRF5; UNIPARC:UPI00000C4DD7; GB:AE002379;
A;Cross-references: UNIPROT:Q9JRF5; UNIPARC:UPI00000C4DD7; GB:AE002379;
A;Cross-references: UNIPROT:Q9JRF5; UNIPARC:UPI0000C4DD7; GB:AE002379;
A;Cross-references: UNIPROT:Q9JRF5; UNIPARC:UPI00000C4DD7; GB:AE002379;
A;Cross-references: UNIPROT:Q9JRF5; UNIPROT:UPI00000C4D07; GB:AE002379;
A;Cross-references: UNIPROT:Q9JRF5; UNIPROT:UPI00000C4D07; GB:AE002379;
A;Cross-references: UNIPROT:UPI0000C4D07; GB:AE002379;
A;Cross-references: UNIPROT:UPI0000C4D07; GB:AE002379;

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A;Authors: Grandi, G.; Sun, L.; Smith, H.O.; Fraser, C.M.; Moxon, E.R.; A;Authors: Grandi, G.; Sun, L.; Smith, H.O.; Fraser, C.M.; Moxon, E.R.;
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A;Accession: T40337
A;Accession: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-450 <LYN>
A;Residues: 1-450 <LYN>
A;Cross-references: UNIPROT:O59711; UNIPARC:UPI0000069880; EMBL:AL022244; PIDN:CAA18292.
A;Experimental source: strain 972h-; cosmid c3B8
C;Genetics:
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A;Map position: 2
C;Keywords: oxidoreductase
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A;Accession: C81996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Title: Complete genome sequence of Neisseria meningitidis serogroup B A;Reference number: A81000; MUID:20175755; PMID:10710307
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A; Residues: 1-504 < PAR>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;Cross-references: UNIPARC:UPI00000C4DD7; GB:AL162752; ;Experimental source: serogroup A, strain Z2491
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;Title: Complete DNA sequence of a serogroup A strain of Neisseria menigitidis Z2491.;Reference number: A81775; MUID:20222556; PMID:10761919
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Pred. No. 3.8e+02;
1; Mismatches 1;
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3.4e+02;
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                        #text_change 09-Jul-2004
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Quail, M.A.;
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; Pizza, M.
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                                                                                                                    A;Molecule type: mRNA
A;Residues: 1-524 <LYN>
A;Cross_references: UNIPROT:P26590; UNIPARC:UPI000012FDFF;
                                                                                                                                                                                                                                                                                                                                                                                                                N;Alternate names: nucleoprotein
C;Species: parainfluenza virus type 1
C;Date: 31-Mar-1992 #sequence_revision 31-Mar-1992 #text_change 09-Jul-2004
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A;Title: The nucleoproteins of human parainfluenza virus
A;Reference number: A38401; MUID:91202138; PMID:1707951
A;Accession: A38401
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R;Lyn, D.; Gill, D.
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                                                                                               Superfamily:
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nucleocapsid;

70.2%;

Score 33;

DB ۳,

Length

paramyxovirus

nucleocapsid

protein

GB:D01070; NID:g222271;

PIDN

D.S.; Scroggs, R.A.; Portner,

type

and

Sendai virus share

parainfluenza virus type

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C;Superfamily: paramyxovi
C;Keywords: nucleocapsid;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nucleocapsid protein - parainfluenza virus type 1 (strains A1426, 86-315, 6 N;Alternate names: nucleoprotein C;Species: parainfluenza virus type 1 C;Date: 17-Feb-1994 #sequence_revision 17-Feb-1994 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Experimental source: strain v26, developing flowers C;Superfamily: petunia myb-related protein 3; myb DNA-binding repeat hor C;Keywords: DNA binding; duplication; nucleus; transcription regulation F;25-77/Domain: myb DNA-binding repeat homology <MYB1> F;78-128/Domain: myb DNA-binding repeat homology <MYB2>
                                                                                                                                                                                                    A;Gene: NP
                                                                                                                                                                                                                                             A;Cross-references: UNIPROT:P36354; UNIPARC:UPI000012FDFD; GA;Note: sequence extracted from NCBI backbone (NCBIN:106078,
                                                                                                                                                                                                                                                                                                                                                                                      R;Miyahara, K.; Kitada, S.; Yoshimoto, M.; Matsumura, H.; Kawano, Arch. Virol. 124, 255-268, 1992
A;Title: Molecular evolution of human paramyxoviruses. Nucleotide
                                                                                                                                                                                                                            C;Genetics:
                                                                                                                                                                                                                                           A; Note: sequence extracted
                                                                                                                                                                                                                                                                                                       A; Molecule type:
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A; Residues: 1-517 < AVI>
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A; Accession: $26606
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                                                                                                                                                                                                                                                                                        A;Residues:
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B, NCBIP:106079)
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R;Kurodá, M.; Ohta, T.; Uchiyama, I.; Baba, T.; Yuzawa, H.; Kobayashi, I.; Cui, L.; ma, A.; Mizutani-Ui, Y.; Kobayashi, N.; Sawano, T.; Inoue, R.; Kaito, C.; Sekimizu, C.; Shiba, T.; Hattori, M.; Ogasawara, N.; Hayashi, H.; Hiramatsu, K. Lancet 357, 1225-1240, 2001

A;Title: Whole genome sequencing of meticillin-resistant Stapylococcus aureus.

A;Reference number: A89758; MUID:21311952; PMID:11418146
              hypothetical protein T6G15.10 - Arabidopsis thaliana
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 23-Apr-1999 #sequence_revision 23-Apr-1999 #text_change 09-Jul-2004
                                                                                RESULT 100
T06648
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A;Reference number: A71570; MUID:99000809; PMID:9784136
A;Accession: C71529
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Stephens, R.S.; Kalman, S.; Lammel, C.J.; Fan, J.; Marathe, Science 282, 754-759, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hypothetical protein CT326 - Chlamydia trachomatis (serotype D, strain UM3/Cx) (Species: Chlamydia trachomatis C;Species: Chlamydia trachomatis C;Date: 13-Sep_1998 #sequence_revision 13-Sep_1998 #text_change 09-Jul-2004 C;Accession: C71529
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A;Status: preliminary
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C;Species: Staphylococcus aureus
C;Date: 10-May-2001 #sequence_revision 10-May-2001 #text_change 09-Jul-2004
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2; Mismatches
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Pred. No. 4.3e+02;
1; Mismatches 1;
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A; Residues: 1-650 <BEV>
                                                                                                                                                                                                                                             A; Reference number: Z15791
A; Accession: T06648
                                                                                                                                                                                                                                                                            R;Bevan, M.; Murphy, G.; Ridley, P.; Hudson, S.; Bancroft, I.; Mewes, H.W.; Mayer, submitted to the Protein Sequence Database, April 1999
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Q59MN9 CANAL
Q81E68 PLAF7
Q51L08 ORYSA
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Q5AB5 CANAL
Q64Q25 CANAL
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Q64Q31 THEPA
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Q97LF8 CLOAMB
Q3YT66 EHRCJ
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Q54U43_DICDI
Q38DT4_9TRYP
Q6C9S5_YARLI
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Q81AM1_PLAFA
Q81ST86_PLAFA
Q65859_PLAFA
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Q7R894_PLACH
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Q5TKF1_ORYSA
Q4Y3M3_PLACH
Q2ZA919_GAMM
Q356F4_9GAMM
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Q8ILD1 PLAF7
PDS5 YEAST
Q6FMZ8 CANGA
Q8I3B1 PLAF7
Q8IBA8 PLAF7
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Q6ZTY1_ORYSA
Q5XX08_9MONO
Q94TC1_DROME
Q21C53_CAEEL
Q8SEP3_NEUCR
Q8SEP3_NEUCR
Q7SC2_NEUCR
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Q21053

O8ssp3

Q7ssp3

Q7ssp3

Q7ssp3

Q7ssp3

Q5m6g1

Q5m6g1

Q5cv1

Q5si6

Q6cv1

Q5y51

Q7x17

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Q4z8k6

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ID SCKA PANIM
STANDARD; PRT;
AC P55927;
AC P55927;
DT 01-NOV-1997, integrated into UniProtK
DT 15-DEC-1998, sequence version 2.
DT 07-FEB-2006, entry version 47.
DE potassium channel-blocking toxin PiTX
DE alpha) (Potassium channel-blocking toxin PiTX
DE potassium channel-blocking toxin PiTX
DE pandinus imperator (Emperor scorpion)
OC Likaryota, Metazoa, Arthropoda; Cheli
OC Inrida; Scorpionoidea; Scorpionidea;
NUCLEOTIDE SEQUENCE (MRNA).
RA ROGOWSKI R.S., Collins J.H., O'Neill
RA Werkman T.R., Rogawski M.A., Tenenhol
RA Werkman T.R., Rogawski M.A.
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Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachmida; Scorpiones; Iurida; Scorpionoidea; Scorpionidae; Scorpioninae; Pandinus.

NCBI_TaxID=55084;

[1]
                                                                                                                                                                                                                               MEDLINE=97070422; PubMed=8913348; Rogowski R.S., Collins J.H., O'Neill T.J., Gustafson Werkman T.R., Rogawski M.A., Tenenholz T.C., Weber D. Blaustein M.P.;
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l. 50:1167-1177(1996).
                                                                                                         OF 13-47, AND FUNCTION
       PubMed=8660410; DOI=10.1007/s002329900084; Olamendi-Portugal T., Zamudio F.Z., Possan: from the venom of the scorpion Pandinus imp
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Q8STF4 PLAFA
Q8A71 BACTM
Q866X1 PLAFA
Q8A71 PLAFA
Q8A71 PLAFA
Q8MSG6 STRT1
Q5MSG6 STRT2
Q59Q2 CANAL
Q77177 PLAFA
Q3BHK9 PLAFA
Q3BHG7 PLAFA
Q3BHG7 PLAFA
Q3BHG7 PLAFA
Q3BHG7 PLAFA
Q3BJG5 PLAFA
Q3BJG7 CANEL
Q3ZY49 GTRSU
Q1739 CABER
Q5ANM5 CANAL
Q3CHO THEET
Q3TIL4 MOUSE
Q3TIL4 MOUSE
Q3TIL4 RATTH
Q816F7 BACCR
Q3ERMO BACTI
Q3GRM0 BACTI
Q75419 DICDI
Q5419 DICDI
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Tenenholz T.C., Weber D.
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Q8aa71
Q86ax1
Q86ax96
Q5m596
Q5m596
Q5m595
Q5m595
Q5phk9
Q3bhk9
Q3bj16
Q3bf51
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Q3bf51
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RESULT
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Matches &
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N6 HUMAN
QBNFN6 HUMAN
QBNFN6;
        Eukaryota; Metazoa;
Mammalia; Eutheria;
                           Homo
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DISULFID
DISULFID
NON TER
STRAND
                                    Name=RGS3;
                                           RGS3 isoform
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TURN
STRAND
                                                                                                                                                                                                                                                                                                                                                                              Pfam; PF00451; Toxin 2; 1.
PRINTS; PR00286; CHARYBDTOXIN
ProDom; PD003586; Scorpion tox
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Copyrighted by the UniProt Consortium, Distributed under the Creative Commons
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   show that the N-terminal amino acid sequence is important for their affinities towards Shaker B K+ channels."; J. Membr. Biol. 152:49-56(1996).
                                                                                                                                                                                                                                    STRAND
                                                                                                                                                                                                                                                                                                                                                      Neurotoxin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=97215813; PubMed=9062103; DOI=10.1021/bi9628432; Tenenholz T.C., Rogowski R.S., Collins J.H., Blaustein M.P.,
                                                                                                                                                                                                                                                                                                                                      PEPTIDE
                                                                                                                                                                                                                                                                                                                                              SIGNAL
                                                                                                                                                                                                                                                                                                                                                             3D-structure; Direct protein
                                                                                                                                                                                                                                                                                                                                                                        PROSITE;
                                                                                                                                                                                                                                                                                                                                                                                                      InterPro; IPR001947; Scorpion_toxinS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Weber D.
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                                                                                                                                                                                                                                                                                                                                                                                                                     LinkHub; P55927; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SUBCELLULAR LOCATION: Secreted pr
TISSUE SPECIFICITY: Expressed by
SIMILARITY: Belongs to the short
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         channel inhibitor subfamily.
                          sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                          2PTA; NMR; @=13-47.
                                                                                                                                                                                                                                                                                                                                                                                                                                    T10471;
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                                                                                                                                            Similarity
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ilarity 100.0%;
Conservative 0
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                                           sequence version entry version 13. RGS3S.
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A
                                                                                      PRELIMINARY;
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SCORP_SHORT_TOXIN; 1.
ect protein sequencing;
sium channel inhibitor;
                                                                                                                                   12
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                                                                                                                                                                                                         5434
      Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Primates; Catarrhini; Homin
                                                                                                                                                                                                                           MW;
                                                                     into UniProtKB/TrEMBL
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                                                                                                                                                                                                                                                                                                                                                                                                                                             mRNA.
                                                                                     PRT;
                                                                                                                                                                            Score 47; I
Pred. No. 2.
                                                                                                                                                                                                        21B8BF110A37BD7C CRC64;
                                                                                                                                                                                                                                                                                                                                   Potassium
                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 the venom
                                                                                      319
                                                                                                                                                                                                                                                                                                                                  channel-blocking
                                                                                                                                                                                                                                                                                                                                                                                                                                                           see http://www.uniprot.org/terms
Attribution-NoDerivs License
                                                                                                                                                                                     DB 1; Length 47;
                                                                                                                                                                                                                                                                                                                                                     Signal;
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                                                                                                                                                                                                                                                                                                                                                            inhibitor;
                                                                                                                                                                                                                                                                                                                                   toxin PiTX-K-
      Hominidae;
                                                                                                                                                                    0;
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RESULT 3

1036046 MA

1036046 MA

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Best Local
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  Signal
SIGNAL
                                     EMBL;
                                                                                                                                                    ORFNames=MaquDRAFT_1112;
Marinobacter aquaeolei VT8.
Bacteria; Proteobacteria; Gammaproteobacteria;
                                                                                                     Copyrighted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-DEC-2005, integrated int
06-DEC-2005, sequence versi
07-FEB-2006, entry version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRINTS; PR01301; RGSPROTEIN.
ProDom; PD001180; Reg1_Gprotein;
SMART; SM00315; RGS; 1
PR0SITE; PS50132; RGS; 1
                                                                                 Distributed
                                                                                                                                                                                                                                                                       US DOE Joint Genome Institute (JGI-ORNL);
                                                                                                                                                                                                                                                                                                    STRAIN=VT8
                                                                                                                                                                                                                                                                                                                     NUCLEOTIDE
                                                                                                                                                                                                                                                                                                                                                               Submitted
                                                                                                                                                                                                                                                                                                                                                                                                 US DOE Joint Genome Institute (JGI-PGF);
COpeland A., Lucas S., Lapidus A., Barry K.,
Hammon N., Israni S., Pitluck S., Richardson
"Sequencing of the draft genome and assembly
                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN=VT8;
US DOE Joint
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUCLEOTIDE SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=351348;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Alteromonadaceae; Marinobacter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Q36Q46_MARHY
Q36Q46;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=22032988; PubMed=12036301; DOI=10.1006/geno.2002.6773; Kehrl J.H., Srikumar D., Harrison K., Wilson G.L., Shi C.S.; "Additional 5' exons in the RGS3 locus generate multiple mRNA transcripts, one of which accounts for the origin of human PD7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Uncharacterized conserved membrane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GO; GO:0004871; F:signal
InterPro; IPR000342; RGS.
Pfam; PF00615; RGS; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; AF490838; AAM33253.1; -; mRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Copyrighted by the UniProt Cons
Distributed under the Creative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       transcripts, one of which accounts for Genomics 79:860-868(2002).
-!- SIMILARITY: Contains 1 RGS domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ensembl; ENSG00000138835;
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                                                                                                                                             preliminary data.
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                                          AALG01000011;
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRELIMINARY;
                                                                                 the Creative
                                                                                                        UniProt
                                       EA098985.1; -;
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                                                                                                        Consortium,
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Pred. No. 17;
                                                                               Sortium, see http://www.uniprot.org/terms Commons Attribution-NoDerivs License
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                                     Genomic_DNA
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Attribution-NoDerivs License
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                                                                                                                                                                                                                                                                                                                                                                                                   Marinobacter
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                                                                                                                                                                   which
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.S.;
                                                                                                                                                                                                                                 VT8.";
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RESULT 5
Q7YZ83 PLACH
ID Q7YZ83, PLACH
AC Q7YZ83;
DT 01-CCT-2003, s
DT 07-FEB-2006, e
DE PC10106C.
GN ORFNames=PC101
OS Plasmodium cha
OC Eukaryota; Alv
OX NCBI TaxID=312
RN [1]
RP NUCLEOTIDE SEC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-JUL-2005, integrated into UniProtKB/TrEMBL.
05-JUL-2005, sequence version 1.
07-FEB-2006, entry version 5.
Hypothetical protein (Fragment).
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PubMed=15637271; DOI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             InterPro; IPR006484; PYST_B.
TIGRFAMS; TIGR01597; PYST-B;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Distributed
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Science
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               transcriptomic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "A comprehensive survey of the Plasmodium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sinden R.S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Plasmodium chabaudi
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                                                                            Plasmodium chabaudi chabaudi.
Eukaryota; Alveolata; Apicomp
                                                                                                                              ORFNames=PC10106c;
     NUCLEOTIDE SEQUENCE
                                               NCBI_TaxID=31271;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PLACH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ence 307:82-86 (2005).

CAUTION: The sequence shown her EMBL/GenBank/DDBJ whole genome
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Lafontaine I., de Montigny J., Marck C., Neuvegilse C., Talla E.,
A. Lafontaine I., de Montigny J., Marck C., Neuvegilse C., Talla E.,
A. Goffard N., Frangeul L., Aigle M., Anthouard V., Babour A.,
Babour A., Babour A.,
Barnay S., Blanchin S., Beckerich J.-M., Beyne E., Bleykasten C.,
A. Boisrame A., Boyer J., Cattolico L., Confanioleri F., de Daruvar A.,
Boisrame A., Boyer J., Cattolico L., Confanioleri F., de Daruvar A.,
A. Despons L., Fabre E., Fairhead C., Ferry Dumazet H., Groppi A.,
A. Hantraye F., Hennequin C., Jauniaux N., Joyet P., Kachouri R.,
A. Kerrest A., Koszul R., Lemaire M., Lesur I., Ma L., Muller H.,
A. Kerrest A., Koszul R., Lemaire M., Lesur I., Ma L., Muller H.,
A. Nicaud J.-M., Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,
A. Nicaud J.-M., Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,
A. Pellenz S., Potier S., Richard G.-F., Straub M.-L., Suleau A.,
A. Pellenz S., Potier S., Wesolowski-Louvel M., Westhof E., Witth B.,
A. Zeniou-Meyer M., Zivanovic Y., Bolotin-Fukuhara M., Thierry A.,
A. Bouchier C., Caudron B., Scarpelli C., Gaillardin C., Weissenbach J.
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Best Local S
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QGFN57;
19-JUL-2004, integrated into U.
19-JUL-2004, sequence version
107-FEB-2006, entry version 16.
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MEDLINE=2672904; PubMed=12707350;
DOI=10.1046/j.1365-2958.2003.03491.x;
Fischer K., Chaychich M., Huestis R., Wilson D.W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBL; AY149028; AAO06133.1; -; InterPro; IPR006484; PYST_B. TIGRFAMS; TIGR01597; PYST-B; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similar to sp|P38748 Saccharomyces cerevisiae YHL010c
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                                                                                                                                                                                                                                              Copyrighted Distributed
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                                                                                                                                                                                                                                                                                                                                       Nature 430:35-44(2004)
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Wincker P., Souciet J.-L.;
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PubMed=15229592; DOI=10.1038/nature02579;
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    InterPro; IPR011422;
InterPro; IPR001841;
InterPro; IPR001607;
                                                                                                                                                                                                                                                                                                                   -!- SIMILARITY: Contains 1 RING-type
                                                                                                                                                                                                                                                                                                                                                             "Genome evolution in yeasts.";
                                                                     L; CR380957; CAG61298.1; -; Genomic_DNA.
G0:0000151; C:ubiquitin ligase complex; IEA.
G0:0046872; F:metal ion binding; IEA.
G0:0004842; F:ubiquitin-protein ligase activity;
G0:0008870; F:zinc ion binding; IEA.
G0:00016567; P:protein ubiquitination; IEA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bleykasten C.,
                                                                                                                              IEA.
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RESULT 7
Q2PEW5 TR
ID Q2PE
DT Q2PE
AC Q2PE
AC Q2PE
AC Q2PE
DT 07-I
DT 07-I
DT 07-I
DT 07-I
OC Spe
OC Tri
OX NCI
RN NCI
RN NCI
RN NCI
RN II6
RT CI
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Q6FK24
ID Q6
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DT 19
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DE S1
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OS Ca
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Best Local S
Matches
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Best Local
                                                                                        Q6FK24 CANGA
Q6FK24;
19-JUL-2004,
19-JUL-2004,
19-FEB-2006,
                        Similar to sp|P35177 Saccl
Transcriptional activator
OrderedLocusNames=CAGLOMO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pfam; PF07576; BRAP2; 1.
Pfam; PF00097; zf-C3HC4; 1.
Pfam; PF00097; zf-C3HC4; 1.
Pfam; PF02148; zf-CUBP; 1.
SMART; SM00184; RING; 1.
SMART; SM00290; ZF UBP; 1.
PROSITE; PS50089; ZF RING 2; 1.
Complete proteome; Metal-Dinding; Zinc; Zinc-finger.
Complete S86 AA; 68323 MW; 550DFA56268A1321 CRC64;
    Candida
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sato S., Isobe S., Asamizu E., Nakamura Y., Ohmido N., S
Klimenko I., Sasamoto S., Wada T., Watanabe A., Tabata S
"Comprehensive structural analysis of the genome of red
(Trifolium gratense).";
Submitted (OCT-2005) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PubMed=13679984; DOI=10.1007/s00122-003-1412-z; Isobe S., Klimenko I., Ivashuta S., Gau M., Kozlov N.N.; Isobe S., Firmenko I., Todover (Trifolium pratense L.) "First RFLP linkage map of red clover (Trifolium pratense L.) cDNA probes and its transferability to other red clover germpi Theor. Appl. Genet. 108:105-112(2003).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Trifolium pratense (Red clover).

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicotyledons; rosids; eurosids I; Fabales; Fabaceae; Papilionoideae; Trifolieae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-FEB-2006, integrated into UniProtKB/TrEMBL.
07-FEB-2006, sequence version 1.
07-FEB-2006, entry version 1.
Putative phosphoinositide specific phospholipase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Distributed under the Creative
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Q2PEW5;
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NCE 603 A
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7; Conserv
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  glabrata
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                                                                                             sequence version entry version 13.
                                                                                                                                         integrated into UniProtKB/TrEMBL
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                                                                                                                                                                                        PRELIMINARY;
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                                                                                                                                                                                                                                                                                                              317
  (Yeast)
                           CAGL0M01738g;
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100.0%;
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Pred. No.
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Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FA71D84631E30A31 CRC64;
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glabrata)
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bata S.;
                                                                    SPT7
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RESULT 9
Q918J4_NPVST
ID Q918J4_NPVST
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Best Local S
Matches 7
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A Lafontaine I., de Montigny J., Marck C., Neuveglise C., Talla E.,
A Goffard N., Frangeul L., Aigle M., Anthouard V., Babour A.,
Barbe
Barnay S., Blanchin S., Beckerich J.-M., Beyne E., Bleykasten C.,
A Boiszame A., Boyer J., Cattolico L., Confanioleri F., de Daruvar A.
Despons L., Fabre E., Fairhead C., Ferry-Dumazet H., Groppi A.,
A Hantraye F., Hemnequin C., Jauniaux N., Joyet P., Kachouri R.,
A Nicaud J.-M., Nikolski M., Oztas Y., Ozier-Kalogeropoulos O.,
A Kerrest A., Koszul R., Lemaire M., Lesur I., Muller H.,
A Kerrest A., Koszul R., Lemaire M., Lesur I., Suleau A.,
A Nicaud J.-M., Nikolski M., Oztas Y., Ozier-Kalogeropoulos O.,
A Pellenz S., Potier S., Richard G.-F., Straub M.-L., Suleau A.,
A Pellenz S., Potier S., Richard G.-F., Straub M.-L., Suleau A.,
A Swennen D., Tekaia F., Wesolowski Louvel M., Westhof E., Wirth B.,
A Swennen D., Tekaia F., Wesolowski Louvel M., Westhof E., Wirth B.,
A Swennen B., Zoudron B., Scarpelli C., Gaillardin C., Weissenbach A.
A Bouchier C., Caudron B., Scarpelli C., Gaillardin C., Weissenbach A.
Mincker P., Souciet J.-L.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRINTS; PR00503; BROMODOMAIN.
SMART; SM00297; BROMO; 1.
PROSITE; PS00633; BROMODOMAIN 1
PROSITE; PS50014; BROMODOMAIN 2
Complete proteome.
SEQUENCE 1385 AA; 157899 MW.
                                                                                                                                                                                                                                                                      01-DEC-2001, integrated into U
01-DEC-2001, sequence version
07-FEB-2006, entry version 10.
Hypothetical protein.
Spodoptera litura multicapsid
                                                                                   "Sequence analysis of the nucleopolyhedrovirus genom Virology 287:391-404(2001)
           EMBL; AF325155; AAL01710
                                         Distributed
                                                      Copyrighted
                                                                                                                                  Yang
                                                                                                                                                                 MEDLINE=21425398;
                                                                                                                                                                                              NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                         Nucleopolyhedrovirus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Distributed under the Creative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Genome evolution in yeasts.";
Nature 430:35-44(2004).
                                                                                                                                                                                                                         NCBI_TaxID=46242;
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NCBI_TaxID=5478;
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PubMed=15229592; DOI=10 1038/nature02579;
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Saccharomycetales; mitosporic
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                                                                                      lyhedrovirus genome."; 287:391-404(2001).
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PS50014; BROMODOMAIN_2;
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                                                                                                                                                Wang
                                                                                                                                             PubMed=11531416; DOI=10.1006/viro.
ang L., Hu X., Bao W., Li G., Chen
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                                                         UniProt
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                                      iProt Consortium, see http://www.uniprot.
Creative Commons Attribution-NoDerivs Li
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                                                                                                                  Spodoptera
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 42; DB
Pred. No. 5.2e
1; Mismatches
                                                                                                                                                                                                                                                                        nucleopolyhedrovirus (SpltMNPV).
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         Genomic_DNA
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                                                                                                                     litura
                                                                                                                                                                                                                                                           Baculoviridae;
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Han H.,
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RA Jacq C., Alt-Moerbe J., Andre B., Arnold W., Bahr A., Ballesta J.P.G., RA Bargues M., Baront L., Becker A., Biteau W., Bloecker H., Blugeon C., RA Boskovic J., Brandt P., Brueckner M., Buitrago M.J., Coster F., RA Boskovic J., Brandt P., Brueckner M., Buitrago M.J., Coster F., RA Boskovic J., Brandt P., Brueckner M., Buitrago M.J., Coster F., RA Boskovic J., Brandt P., Legros Y., Louis E.J., Carcia-Cantalejo J.M., RRA Goffeau A., Gomez-Peris A., Granotier C., Hanemann V., Hankeln T., RA Goffeau A., Gomez-Peris A., Granotier C., Hanemann V., Hankeln T., RA Hoheisel J.D., Jaeger W., Jimenez A., Joniaux J.-L., Kraemer C., RA Monnet A., Moro M., Perez-Alonso M., Perez-Ortin J.E., Pohl T.M., RA Rieger M., Salom D., Saluz H.P., Saiz J.B., Saren A.-M., Schwarz S., Narpid H., Purnelle B., Rasmussen S.W., Remacha M.A., Revuelta J.L., RA Rieger M., Salom D., Saluz H.P., Saiz J.B., Saren A.-M., Schwarz S., Voltkaert G., Wagner G., Wambutt R., Wedler E., Wedler H., Woelfil S., RA Voltkaert G., Wagner G., Wambutt R., Wedler E., Medler H., Woelfil S., RA Niblett D., Odell C., Oliver K., Rajandream M.A., Richards C., RA Harris D.E., Bomman S., Brown D., Churcher C.M., Connor R., Dedman K., RA Harris D.E., Araujo R., Aviles B., Berno A., Carpenter J., Chen B., Cherry J.M., Chung E., Duncan M., Hunicke-Smith S., Hyman R.W., RA Namath A., Oefner P., Oh C., Petel F.X., Roberts D., Schramm S., RA Namath A., Oefner P., Oh C., Petel F.X., Roberts D., Schramm S., RA Namath A., Oefner T., Shroff N., Winant A., Yelton M.A., RA Cooper J., Blaskari D., Lew H., Lin D., Mosedale D., Nakahara K., RA Greco T., Hallsworth K., Hawkins J., Hiller L.W., Jier M., RA Cooper J., Blaskari D., Ratten J., Kucaba T., Langston Y., RA Latreille P., Le T., Mardis E., Menezes S., Miller N., Nhan M., Ra Kerston R., Albermann K., Hani J., Heumann K., Kleine K., Mewes H.-W., Zollner A., Zaccaria P., Waddin M., Wilson R.W., Tevaskkis E., Racharomyces of erryisise chromosome IV.".
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Q04921;
Q04921;
126-APR-2004, integrated into U
01-NOV-1996, sequence version 3.
07-MAR-2006, entry version 33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Saccharomyces cerevisiae (Baker's yeast).

Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
                    Microbiology 142:2897-2905(1996).
-!- FUNCTION: Septins are GTPases involved in cytokinesis that assemble into filaments and form a ring at the cleavage si
                                                             de Virgilio C., DeMarini D.J., Pringle J.R "SPR28, a sixth member of the septin gene cerevisiae that is expressed specifically
                                                                                                                                                                       Mewes H.-W., Zollner A., "The nucleotide sequence Nature 387:75-78(1997).
                                                                                                                                         SUBCELLULAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=97313263; PubMed=9169867;
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HOF1, a
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a protein
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    the cleavage site. involved in septati
                                                                                                                                         INTERACTIONS WITH CDC11
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RESULT 11
Q2IWZ5 RHOPA
Q2IWZ5 RHOPA
PRELIMINARY;
AC Q2IWZ5;
AC Q2IWZ5;
DT 07-MAR-2006, integrated into
DT 07-MAR-2006, sequence version
DT 07-MAR-2006, entry version 1.
DE Trigger factor.
GN ORFNames=RPB 2562;
GN CAFNames=RPB 2562;
OS Rhodopseudomonas palustris Ha
OC Bacteria; Proteobacteria; Alg
OC Bradyrhizobiaceae; Rhodopseud
OX NCBI_TaxID=316058;
RN [1]
RP NUCLEOTIDE SEQUENCE.
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Best Local
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COTLED
COMPBIAS
                                                               ORFNames=RPB_2562;
Rhodopseudomonas palustris HaA2.
Bacteria; Proteobacteria; Alphaproteobacteria; Rhizobiales;
                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
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sequence version

Rhodopseudomonas

integrated into UniProtKB/TrEMBL

PRT;

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Linkhub; Q04921;
GO; GO:0000144; C
GO; GO:0005515; F
GO; GO:0005200; F
GO; GO:0007047; P
GO; GO:0009902; P
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SGD; S00002626; SPR28.
BioCyc; SCRP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                         Promom; Provision, GTP Cell Div; 1.

Probom; PD002565; GTP Cell Div; 1.

Cell cycle; Cell division; Coiled coil; Con
Cell cycle; Cell division; Nucleotide-binding;
GTP-binding; Membrane; Nucleotide-binding-ri
CHAIN 1 423
                                                                                                                                                                                                                                                                                                                                                                        InterPro; IPR000038; Cell Div GTP PANTHER; PTHR18884; GTP Cell Div; Pfam; PF00735; GTP CDC; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                        GO:0000144; C:septin ring (sensu Saccharomyces); TAS. GO:0005515; F:protein binding; IPI. GO:0005500; F:structural constituent of cytoskeleton; TAS. GO:0005200; F:scell wall organization and biogenesis; TAS. GO:0000902; P:cellular morphogenesis; TAS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   formation. First expressed at the beginning highly expressed prior meiosis II. SIMILARITY: Belongs to the septin family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 P32458:CDC11; NDExp=2; IntAct=EBI-38416, EBI-4178; SUBCELULLAR LOCATION: Membrane-associated. Present at the bud r during cell division. Probably interacts with phosphoinosides as phosphatidylinositol 4-phosphate or phosphatidylinositol 5-phosphate (By similarity). Localizes to ring-like structures around each of the four nuclear lobes at the onset and during meiosis II. Concentrated initially at the leading edge of the developing prospore wall.

DEVELOPMENTAL STAGE: Expressed during meiosis and ascospore between the process of the developmental stages around the developmental stages around the developmental stages around the developmental stages around the developmental stages are developmental stages are developmental stages.
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SUBUNIT: Interacts with
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384
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423 AA;
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                                                                                                       Score 41; DB 1;
Pred. No. 2.2e+02;
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EB8D3C0F47CD41DB CRC64;
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Potential.
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QBJKJ5;
01-OCT-2002,
01-OCT-2002,
07-FEB-2006,
Orf118.
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05-JUL-2004,
07-FEB-2006,
                                                                                                                                                                                                                                                                                                                       STRAIN=ATCC VR-111;
PubMed=14671117; DOI=10.1128/JVI.78.1.353-366.2004;
PubMed=14671117; DOI=10.1128/JVI.78.1.353-366.2004;
Tulman E.R., Afonso C.L., Lu Z., Zsak L., Kutish G.F.,
"The genome of canarypox virus.";
J. Virol. 78:353-366(2004).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Copeland A., Lucas S., Lapidus A., Barry K., Detter J.C., Glavina Hammon N., Israni S., Pitluck S., Chain P., Malfatti S., Shin M., Vergez L., Schmutz J., Larimer F., Land M., Hauser L., Pelletier Kyrpides N., Anderson I., Oda Y., Harwood C.S., Richardson P.; "Complete sequence of Rhodopseudomonas palustris HaA2."; Submitted (JAN-2006) to the EMBL/GenBank/DDBJ databases.
                       Name=orf118;
                                                                                                                                                                                                                                                                                    Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=44088;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Canarypox virus (CNPV).
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Q6VZM2;
           Heliothis
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9VIRU
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NCE 1767 AA; 199286
                                                                                                                                                                                                                                                    AY318871; AAR83471.1; -; Genomic_DNA.
Pro; IPR007490; Poxvirus_B22R.
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VCE 452 AA; 50111 MW; 4F805CDC1AC12E5D
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                                                                                              PubMed=15994558; DOI=10.1126/science.1110439;
PubMed=15994558; DOI=10.1126/science.1110439;
Gardner M.J., Bishop R., Shah T., de Villiers E.P., Carlton J.M.,
Hall N., Ren Q., Paulsen I.T., Pain A., Berriman M., Wilson R.J.,
Sato S., Ralph S.A., Mann D.J., Xiong Z., Shallom S.J., Weidman J.,
Jiang L., Lynn J., Weaver B., Shoalid A., Domingo A.R., Wasawo D.,
Crabtree J., Wortman J.R., Haas B., Angiuoli S.V., Creasy T.H., Lu (
Suh B., Silva J.C., Utterback T.R., Feldblyum T.V., Pertea M.,
Allen J., Nierman W.C., Taracha E.L., Salberg S.L., White O.R.,
Fitzhugh H.A., Morzaria S., Venter J.C., Fraser C.M., Nene V.;
"Genome Sequence of Theileria parva, a Bovine Pathogen That Transfo
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Cheng C.H., Heiao Y.Y.,
Submitted (JUN-2002) to
                                                                                        Science
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DOI=10.1128/JVI.76.18.9024-9034.2002;
Cheng C.H., Liu S.M., Chow T.Y., Hsiao
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NCE 322 AA; 36706 M
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W; 22B25C1209717A90
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InterPro; IPRO01650; Helicase_C.
Pfam; PF00270; DEAD; 1.
Pfam; PF00271; Helicase_C; 1.
SMART; SM00487; DEXDC; 1.
SMART; SM00490; HELICC; 1.
ATP-binding; Helicase; Hydrolase.
SEQUENCE 432 AA; 49387 MW; 02F0BE597876EE1E CRC64;
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EMBL; BX294143; CAD74545.1; -; Genomic DNA GO; GO:0003992; F:acetylornithine transamic GO; GO:0030170; F:pyridoxal phosphate hind:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bacteria; Planctomycetes; Planctomycetacia; Planctomycetaceae; Pirellula.
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01-OCT-2003,
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                                                                                                                                                                                                                        strain 1
                                                                                                                                                                                                                                                                                                                                                                                     NUCLEOTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                            Planctomycetaceae;
NCBI_TaxID=117;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rhodopirellula baltica.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OrderedLocusNames=RB5984;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Probable acetylornithine aminotransferase (EC 2.6.1.11).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              -
                                                                                                                                                                                                                                             "Complete genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GO:0005524; F:ATP binding; IEA.
GO:0008026; F:ATP-dependent helicase a
GO:0016787; F:hydrolase activity; IEA.
GO:0003676; F:nucleic acid binding; IEA.
GO:0000166; F:nucleotide binding; IEA.
                                                                                                                                                                                 SIMILARITY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             preliminary data.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        264 NYKDDDDR 271
                                                                                                                                                                                            n 1.";
Natl. Acad. Sci. U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAGK01000001; EAN34161.1; -;
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6; Conserv
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                                                                                        under the Creative Commons
                                                                                                             by the UniProt Consortium,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRELIMINARY;
                                                                                                                                                                                 Belongs
F:acetylornithine transaminase F:pyridoxal phosphate binding;
                                                                                                                                                                                                                                                sequence of
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                                                                                                                                                                                 the class-III
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                                                                                                                                                                                                                                                the marine planctomycete
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Attribution-NoDerivs License
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Q6C1V6;
22-NOV-2005,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dujon B., Sherman D., Fischer G., Durrens P., Casaregola S.,
Lafontaine I., de Montigny J., Marck C., Neuveglise C., Talla E.,
Goffard N., Frangeul L., Aigle M., Anthouard V., Babour A., Barbe
Barnay S., Blanchin S., Beckerich J.-M., Beyne E., Bleykasten C.,
Bolsrame A., Boyer J., Cattolico L., Confanioleri F., de Daruvar A.
Despons L., Fabre E., Fairhead C., Ferry-Dunazet H., Groppi A.,
Hantraye F., Hennequin C., Jauniaux N., Joyune P., Kachouri R.,
Kerrest A., Koszul R., Lemaire M., Lesur I., Ma L., Miller H.,
Nicaud J.-M., Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,
Pellenz S., Potier S., Richard G.-F., Straub M.-L., Suleau A.,
Swennen D., Tekaia F., Wesolowski-Louvel M., Westhof E., Wirth B.,
Zenjou-Meyer M., Zivanovic V., Polotin-Fubihara M., Miller H.,
Zenjou-Meyer M., Zivanovic V., Polotin-Fubihara M.,
Zenjou-Meyer M., Zivanovic V., Polotin B.,
Zenjou-Meye
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUCLEOTIDE SEQUENCE [LARGE STRAIN=CLIB 122 / E 150;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Saccharomycetales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Yarrowia lipolytica (Candida lipolytica).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRAIN=CLIB 122 / E 150;
PubMed=15229592; DOI=10.1038/nature02579;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-MAR-2006, entry version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GO; GO:0016740;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FUNCTION: Involved in pre-mRNA splicing (By similarity).
SUBUNIT: Associated with the spliceosome (By similarity)
SUBCELLULAR LOCATION: Nucleus (By similarity).
SIMILARITY: Belongs to the CWC25 family.
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                                                                                                                                                                                                                                                                                                                               coil;
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Similarity 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               integrated into UniProtKB/Swiss-Prot
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/FTId=PRO_0000079595
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24-MAY-2005,
07-FEB-2006,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pfam; PF02386; TrkH; 1.
PROSITE; PS00120; LIPASE_SER; UNKNOWN_1.
PROSITE; PS00012; PHOSPHOPANTETHEINE; UNKNOWN_1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hypothetical protein. SEQUENCE 648 AA; 73510 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sucgang R., Berriman M., Song J., Olsen R., Szafranski K., Xu Q Tunggal B., Kummerfeld S., Madera M., Konfortov B.A., Rivero F. Bankier A.T., Lehmann R., Hamlin N., Davies R., Gaudet P., Fey Pilcher K., Chen G., Saunders D., Sodergren E.J., Davis P., Kerhornou A., Nie X., Hall N., Anjard C., Hemphill L., Bason N. Ferrbrother P., Desany B., Just E., Morio T., Rost R., Churcher Character P., Desany B., Just E., Morio T., Rost R., Churcher Character P., Desany B., Just E., Morio T., Rost R., Churcher Character P., Desany B., Just E., Morio T., Rost R., Churcher Character P., Desany B., Just E., Morio T., Rost R., Churcher Character P., Desany B., Just E., Morio T., Rost R., Churcher Character P., Desany B., Just E., Morio T., Rost R., Churcher Character P., Peterbrother P., Peterbrother P., Desany B., Just E., Morio T., Rost R., Churcher Character P., Peterbrother P., Peterbrother
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GO; GO:0008324; F:cation transporter GO; GO:0006812; P:cation transport; I InterPro; IPR008262; Lipase AS. InterPro; IPR006162; Ppantne S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Copyrighted Distributed
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Bichinger L., Pachebat J.A., Gloeckner G., Rajandream M.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hypothetical protein. ORFNames=DDB0184366;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       preliminary data.
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llarity 75.0%;
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                                               integrated into UniProtKB/TrEMBL
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Pred. No. 7.1e+02;
2; Mismatches 0;
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SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=AX4;
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PANTHER; PTHR11070; UvrD-helicase; 1.
Pfam; PF00580; UvrD-helicase; 2.
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Proc. Natl. Acad. Sci. U.S.A.
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Shimizu T., Ohtani K., Hirakawa H., Ohshima K., Yamashita A.,
Shiba T., Ogasawara N., Hattori M., Kuhara S., Hayashi H.;
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MEDLINE=21664373; P
Shimizu T., Ohtani
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Racteria; Firmicutes; Clostridia;
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GO:0004003; F:ATP-dependent DNA
GO:0003677; F:DNA binding; IEA.
GO:0005281; P:DNA repair; IEA.
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QSCQA2;
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12-APR-2005, integrated into
12-APR-2005, sequence versic
07-FEB-2006, entry version 5
Hypothetical protein.
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13-SEP-2005, sequence version
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Probable signal peptide protei
OrderedLocusNames=Reut_A2446,
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-I- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
Copeland A., Lucas S., Lapidus A., Barry K., Detter J.C., Glavina '
Hammon N., Israni S., Pitluck S., Goltsman E., Martinez M.,
Schmutz J., Larimer F., Land M., Lykidis A., Richardson P.,
"Complete sequence of chromosome 1 of Ralstonia eutropha JMP134.",
Submitted (AUG-2005) to the EMBL/GenBank/DDBJ databases.
                          ORFNames=cgd4_1500;
Cryptosporidium parvum
Eukaryota; Alveolata; i
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narreria: Proteobacteria: Betaproteobacteria; Burkholderiales;
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STRAIN=Iowa type II;
STRAIN=Iowa type II;
PubMed=15044751; DOI=10.1126/science.1094786;
PubMed=15044751; DOI=10.1126/science.1094786;
Abrahamsen M.S., Templeton T.J., Enomoto S., Abrahante J.E.,
Abrahamsen M.S., Templeton T.J., Widmer G., Tzipori S., Buck G.J.
Lancto C.A., Deng M., Liu C., Widmer G., Tzipori S., Buck G.J.
Bankler A.T., Dear P.H., Konfortov B.A., Spriggs H.F., Iyer J.
Anantharaman V., Aravind L., Kapur V.,
Anantharaman V., Aravind L., Kapur V.;
"---late Genome sequence of the apicomplexan, Cryptosporidi
                                                                                                                                                                                                                                                                                                                            OrderediocusNames=At5955650;
Arabidopsis thaliana (Mouse-ear cress).
Bukaryota; Viridiplantae; Streptophyta; Embryoph
Spermatophyta; Magnoliophyta; eudicotyledons; cc
rosids; eurosids II; Brassicales; Brassicaceae;
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Sequence features of the regions of 1,456,315
physically assigned Pl and TAC clones.";
DNA Res. 5:41-54(1998).
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MEDLINE=98290546; PubMed=9628582; DOI=10.1093/dnares/5.1.41;
Sato S., Kaneko T., Kotani H., Nakamura Y., Asamizu E., Miya
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Tabata S.;
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Science 304:441-445(2004).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14670 MW;
                                                                                                      17650
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             80.9%;
85.7%;
                                                                80.9%;
85.7%;
                                                                                                      MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    shown here is
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score
Pred.
                                                                Score 38;
Pred. No.
                                                   Pred. No. 2.20
L; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRT;
                                                                                                      7A4C895328B7E2DC
                                                                                                                              Genomic_DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   77D0BDCDFB44094A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        shotgun
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         derived
gun (WGS)
                                                                                                                                                                                                                                                                                                                               Embryophyta; Tracheophyta;
edons; core eudicotyledons;
icaceae; Arabidopsis.
                                                                             DB 2;
                                                                .2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .8e+02;
                                                                                                                                                                                                                                                                                                                                                                                               ū
                                                                                                                                                                                                                                        chromosome
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 125;
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                                                                             Length 154;
                                                                                                       CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                P1 clone:MDF20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       from
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RESULT 24
076542 DICDI
ID 076542 DICDI
AC 076542;
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ID Q8D4G1 V
AC Q8D4C1 V
AC Q8D4C1 V
AC Q8D4C1 V
AC Bacteria OC Vibriona
OC Vibriona
OC Vibriona
OC Vibriona
OC Vibriona
AC VIBRIONA
AC COMPLET
RC STRAINGCA
RA Rhee J.H
RA Choy H.E
RT "Complete
CC -----
CC -----
CC EMBL; BEOCYC; AE
DR BioCyc; AE
DR BIOCY
             S
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RX PubMed=18875012; DOI=10.1038/nature03481;
RA Eichinger L., Pachebat J.A., Gloeckner G., Rajandream M.A.,
RA Sucgang R., Berriman M., Song J., Olsen R., Szafranski K., Xu Q.,
RA Sucgang R., Berriman M., Song J., Olsen R., Szafranski K., Xu Q.,
RA Sungang B., Kummerfeld S., Madera M., Konfortov B.A., Rivero F.,
RA Bankier A.T., Lehmann R., Hamlin N., Davies R., Gaudet P., Fey P.,
RA Balkier A.T., Lehmann R., Hamlin N., Davies R., Gaudet P., Fey P.,
RA Pilcher K., Chen G., Saunders D., Sodergren E.J., Davis P.,
RA Kerhornou A., Nie X., Hall N., Anjard C., Hemphill L., Bason N.,
RA Kerhornou A., Nie X., Hall N., Anjard C., Hemphill L., Bason N.,
RA Farbrother P., Desany B., Just E., Morio T., Rost R., Churcher C.M.,
RA Farbrother P., Desany B., Just E., Morio T., Rost R., Goodhead I.,
RA Farbrother P., Desany B., Just E., Morio T., Rost R., Indasay R.,
RA Muzny D.M., Mourier T., Pain A., Lu M., Harper D., Lindsay R.,
RA Muzny D.M., Wourier T., Pain A., Lu M., Harper D., Lindsay R.,
RA Hauser H., James K.D., Quiles M., Madan Babu M., Saito T.,
RA Mights A., Loulseged H., Mungall K.L., Oliver K., Price C.,
RA Guall M.A., Urushihara H., Hernandez J., Rabbinowitsch E., Steffen D.,
RA Sanders M., Ma J., Kohara Y., Sharp S., Simmonds M.N., Spiegler S.,
RA Tanaka Y., Shaulsky G., Schleicher M., Weinstock G.M., Rosenthal A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-MAR-2003,
07-FEB-2006,
Hypothetical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Complete
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUCLEOTIDE SEQUENCE.

Iranfar N., Loomis W.F.;

Submitted (JUL-1998) to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-NOV-1998, integrated into U
01-NOV-1998, sequence version
21-FEB-2006, entry version 36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Complete
Submitted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRAIN-AX4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ubiquitin-conjugating enzyme pr
Name=ubcC; ORFNames=DDB0219926;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rhee J.H., Kim S.Y., Choy H.E.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OrderedLocusNames=VV21340; ORFNames=VV2_1340; Vibrio vulnificus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          QBD4G1 VIBVU
Q8D4G1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Mycetozoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Dictyostelium discoideum (Slime mold).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bacteria; Proteobacteria;
Vibrionaceae; Vibrio.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=44689;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=CMCP6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AE016796; AAO08227.1; -; Genomic_DNA.

C; VVUL216895:VV21340-MONOMER; -.

ete proteome; Hypothetical protein.

NCE 171 AA; 20363 MW; 985FABC9C2C0)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     integrated into UniProtKB/TrEMBL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRELIMINARY;
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75.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chung S.S.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; Score 38; DB
; Pred. No. 2.5e
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL/GenBank/DDBJ
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985FABC9C2C0A875 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kim J.J., Moon
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RESULT 25
QBL7V6 ARATH
ID QBL7V6, ARATH
AC QBL7V6,
DT 01-CCT-2002, i
DT 07-FEB-2006, e
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Matches 6
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Matches
                                                                                                                                 Query Match
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NUCLEOTIDE SEQUENCE.

Shinn P., Chen H., Cheuk R., Kim C.J., Banh J., Bowser L., Chrinnei P., Chang E., Dale J.M., Goldsmith A.D., Hayashizaki Carninci P., Chang E., Dale J.M., Karlin-Neumann G., Kawai J., L Ishida J., Jones T., Kamiya A., Karlin-Neumann G., Kawai J., L Lee J.M., Lin J., Miranda M., Narusaka M., Nguyen M., Onodera Palm C.J., Quach H.L., Sakurai T., Satou M., Seki M., Southwic Tang C.C., Toriumi M., Wu H.C., Yamada K., Yamamura Y., Yu G., Shinozaki K., Davis R.W., Theologis A., Ecker J.R.;

Shinozaki K., Davis R.W., Theologis A., Ecker J.R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ligase; U
SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DictyBase; DDB0219926; ubcC.
GO; GO:0004842; F:ubiquitin-protein ligase
GO; GO:0006512; P:ubiquitin cycle; IEA.
InterPro; IPR000608; UBQ-conjugat_E2.
Pfam; PF00179; UQ con; I.
SMART; SM00212; UBCc; 1.
                                                                                                                                                                                                         EMBL; AY125517; AAM78108.1; -;
NON_TER 242 242
                                                                                                                                                                                                                                                                           Distributed under the Creative Commons
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NCBI_TaxID=3702;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Arabidopsis thaliana (Mouse-ear cress).

Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicotyledons;
rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PROSITE; PS00183; UBIQUITIN CONJUGAT 1; PROSITE; PS50127; UBIQUITIN CONJUGAT 2;
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Kay R.R., Williams
Kuspa A.;
                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                 Copyrighted
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HSSP; P34477; 1PZV.
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E 235 AA; 27224 MW; F4914E8967359700
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218
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75.0%;
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1; Mismatches
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Pred. No. 3.6e+02;
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3.5e+02;
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, Onodera C.S.,
, Southwick A.,
Y., Yu G., Yu S.,
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NON_TER 2
SEQUENCE 26
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Hall N., Karras M., Raine J.D., Carlton J.M., Kooij T.W.A
Berriman M., Florens L., Janssen C.S., Pain A., Christoph
James K., Rutherford K., Harris B., Harris D., Churcher C
Quail M.A., Ormond D., Doggett J., Trueman H.E., Mendoza
Bidwell S.L., Rajandream M.A., Carucci D.J., Yates J.R. 1
Kafatos F.C., Janse C.J., Barrell B.G., Turner C.M.R., Wa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-FEB-2006, entry vo
Hypothetical protein
ORFNames=PC405060.00
                                                                                                                                                                                           Arabidopsis thaliana (Mouse-ear cress).

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicotyledons;
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Eukaryota; Alveolata; Apicomplexa;
                                                                           Sequence features and BAC clones.";
                                                                                                               MEDLINE=20363099; Put
Kaneko T., Katoh T.,
                                                                                                                                                                                                                                                         01-OCT-2000, sequence version 07-FEB-2006, entry version 15.
                                                                                                                                                                                                                                                                                  Q9LJV8;
01-OCT-2000,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Copyrighted by the UniProt Consortium, Distributed under the Creative Commons
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        transcriptomic, and proteomic analyses."; Science 307:82-86(2005).
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PubMed=15637271; DOI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-JUL-2005, integrated into t
05-JUL-2005, sequence version
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            Kaneko T., Kato T., Sato S., Nakamura Y., Asamizu E., T. Submitted (JUL-1999) to the EMBL/GenBank/DDBJ databases
                                                                                                                                         NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                 OrderedLocusNames=At3g29075;
                                                                                                                                                                                                                                                                                                            Q9LJV8_ARATH
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                                      NUCLEOTIDE
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ARATH
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  preliminary data.
                                                                                                                                                                                                                                                                                                                                                                             103
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                                                                7:217-221 (2000)
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                                                                                                                                                                                                                                                                                                             PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                     7
                                                                                       PubMed=10907853; DOI=10.1093/dnares/7.3.217; I., Sato S., Nakamura Y., Asamizu E., Tabata S.; sis of Arabidopsis thaliana chromosome 3. II. of the 4,251,695 bp regions covered by 90 P1, T2
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85.7%;
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Pred. No.
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                                                                                                                                                                               Brassicaceae;
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Attribution-NoDerivs License
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Churcher C.M.,
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RA Eichinger L., Pachebat J.A., Gloeckner G., Rajandream M.A.,
Sucgang R., Berriman M., Song J., Olsen R., Szafranski K., Xu Q.,
Xa Sucgang R., Berriman M., Song J., Olsen R., Szafranski K., Xu Q.,
Xa Tunggal B., Kummerfeld S., Madera M., Konfortov B.A., Rivero F.,
Xa Pilcher K., Chen G., Saunders D., Sodergren E.J., Davis P.,
Xa Pilcher K., Chen G., Saunders D., Sodergren E.J., Davis P.,
Xa Pilcher P., Desany B., Just E., Morio T., Rost R., Churcher C.M.,
Xa Farbrother P., Desany B., Just E., Morio T., Rost R., Churcher C.M.,
Xa Cooper J., Haydock S., van Driessche N., Cronin A., Goodhead I.,
Xa Munny D.M., Mourier T., Pain A., Lu M., Harper D., Lindsay R.,
Xa Munny D.M., Mourier T., Pain A., Lu M., Harper D., Lindsay R.,
Xa Munny D.M., Mourier T., Pain A., Lu M., Harper D., Lindsay R.,
Xa Mauser H., James K.D., Quiles M., Madan Babu M., Saito T.,
Xa Knights A., Loulseged H., Mungall K.L., Oliver K., Price C.,
Xa Knights A., Urushihara H., Hernandez J., Rabbinowitsch E., Steffen D.
Xa Sanders M., Ma J., Kohara Y., Sharp S., Simmonds M.N., Spiegler S.,
Xa Tivey A., Sugano S., White B., Walker D., Woodward J.R., Winckler T.,
Xa Cox E.C., Chisholm R.L., Gibbs R.A., Loomis W.F., Platzer M.,
Xa Cox E.C., Chisholm R.L., Gibbs R.A., Loomis W.F., Platzer M.
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TAIR;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-FEB-2006, entry version Hypothetical protein. ORFNames=DDB0205379;
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Q54YA8;
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Distributed under the Creative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cox E.C., Chisholm R.L.,
Kay R.R., Williams J.G.,
                                                                                                                                                                                                       Hypothetical p
SEQUENCE 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nature
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                                                                                                                                                                                                                                                                                                                                                                              preliminary data.
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; AB018121; E
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                                                                                                Similarity 6; Conserv
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  DYSDDDDDE
                                                 DYKDDDDK
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nilarity 75.0%;
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                                                                                                   Conservative
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     119
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                                                                                                                                                                                                       35014
                                                                                                                           80.9%;
75.0%;
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Dictyosteliida; Dictyostelium.
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                                                                                                                                                                                                       MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  amoeba Dictyostelium discoideum.";
                                                                                                                                                                                                                                                                                             Consortium, see http://www.uniprot.org/te
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Pred. No. 4.4e
1; Mismatches
                                                                                         Pred. No. 4.00
1; Mismatches
                                                                                                                             Score 38;
Pred. No.
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                                                                                                                                                                                                                                                                                                                               see http://www.uniprot.org/terms
                                                                                                                             .6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                           (WGS)
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                                                                                                                                                                                                         CRC64;
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VHCG_METVO
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Best Local
                                                                                                                  QSAF51 CANAL PRELIMINARY; PRT; QSAF51; Q5AF51; Q5AF51; Q5AF51; Q5AF51; Q5AF72005, integrated into UniPro 26-APR-2005, sequence version 1. 07-FEB-2006, entry version 8. Hypothetical protein. QRFNames=Ca019.10271, Ca019.2757; QRFNames=Ca019.10271, Ca019.2757;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         VHCG METVO
Q00406;
16-JAN-2004,
01-NOV-1996,
07-FEB-2006,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN=DSM 1537 / PS;

STRAIN=DSM 1537 / PS;

PubMed=s121392; DOI=10.1007/BF00281785;

Berghoefer Y., Agha-Amiri K., Klein A.;

Berghoefer Y., Agha-Amiri K., Klein A.;

"Selenium is involved in the negative regulation of the expression selenium-free [NiFe] hydrogenases in Methanococcus voltae.";

Mol. Gen. Genet. 242:369-373(1994).

1- SUBUNIT: The F420-non-reducing hydrogenase vhc is composed of three subunits; vhcA, vhcD and vhcG (By similarity).

1- INDUCTION: By selenium deprivation.

1- SIMILARITY: Belongs to the [NiFe]/[NiFeSe] hydrogenase small subunit family.
  Eukaryota; Fungi; Ascomycota;
Saccharomycetales; mitosporic
NCBI_TaxID=237561;
[1]
                                                                                              Candida albicans SC5314.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
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Distributed
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STRAIN-DSM 1537 / PS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oxidoreductase.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pfam; PF01058; Oxidored_q6;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=92293118;
Halboth S., Klein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Methanococcus voltae.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   InterPro; IPR006137; Oxidored_q6
Pfam; PF01058; Oxidored_q6; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=2188;
                                                                                                                                                                                                                                                                                                CANAL
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S32834; S32834.
                                                                                                                                                                                                                                                                                                                                                                                                 260
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                        integrated into UniProtKB/TrEMBL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             33636 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 38; DB 1;
Pred. No. 4.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           F420-non-reducing G.
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2A88AC3EAE194D8B CF
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                                                Saccharomycotina; Saccharomycetes; Saccharomycetales; Candida.
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Best Local S
Matches 6
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PLACH
Q4XYM1_PLACH
Q4XYM1;
05-JUL-2005,
05-JUL-2005,
07-FEB-2006,
Hypothetical I
                                                                                                                                                                                                     NUCLEOTIDE SEQUENCE.

PubMed=15637271; DOI=10.1126/science.1103717;

Hall N., Karras M., Raine J.D., Carlton J.M., Kooij T.W.A.,

Hall N., Karras M., Raine J.D., Carlton J.M., Kooij T.W.A.,

Hall N., Karras M., Raine J.D., Carlton J.M., Christophides

Berriman M., Florens L., Janssen C.S., Pain A., Churcher C.M.,

James K., Rutherford K., Harris B., Harris D., Churcher C.M.,

Quail M.A., Ormond D., Doggett J., Trueman H.E., Mendoza J.,

Bidwell S.L., Rajandream M.A., Carucci D.J., Yates J.R. III,

Kafatos F.C., Janse C.J., Barrell B.G., Turner C.M.R., Waters
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GO; GO:0005489; F:electron transporter actīvity; GO; GO:000618; P:electron transport; IEA. InterPro; IPR012336; Thiordan-11ke_fd. InterPro; IPR006662; Thiored. InterPro; IPR006662; Thiored. InterPro; IPR006663; Thioredox dom2. InterPro; IPR006663; Thioredox dom2. InterPro; IPR006663; Thioredoxin_fold. Pfam; PF00085; Thioredoxin; 1. PR01235; PR00421; THIOREDOXIN.
                                                                                                                         transcriptomic, and proteomic ar
Science 307:82-86 (2005).
-i- CAUTION: The sequence shown
EMBL/GenBank/DDBJ whole genc
                                                                                                                                                                                                                                                                                                                                                        Plasmodium chabaudi.
Eukaryota; Alveolata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA]
STRAIN=SC5314;
PubMed=15123810; DOI=10.1073/pnas.0401648101;
Jones T., Federspiel N.A., Chibana H., Dungan
             InterPro; IPR000436; S
Pfam; PF00084; Sushi;
                       EMBL; CAAJ01002240; CAH77989.1; -; Genomic_DNA
InterPro; IPR000436; Sushi_SCR_CCP.
                                                                     Copyrighted by the UniProt Consortium, Distributed under the Creative Commons
                                                                                                                                                                   "A comprehensive survey of the Plasmodium transcriptomic, and proteomic analyses.";
                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=5825;
                                                                                                                                                                                                                                                                                                                                                                                     Hypothetical protein (F) ORFNames=PC000687.02.0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hypothetical SEQUENCE 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Copyrighted by the UniProt Consortium, see http://www.uniprot.org/to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "The diploid genome sequence of Candida albicans.";
Proc. Natl. Acad. Sci. U.S.A. 101:7329-7334 (2004).
-!- CAUTION: The sequence shown here is derived from
EMBL/GenBank/DDBJ whole genome shotgun (WGS) ent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Jones T., Federspiel
Magee B.B., Newport G
Davis R.W., Scherer S
                                                                                                                                                                                                   Sinden R.S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EMBL; AACQ01000025;
                                                                                                               preliminary data.
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332 AA; 3
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Newport G., Thorstenson
Scherer S.;
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Attribution-NoDerivs License
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(1)
NUCLEOTIDE SEQUENCE.
STRAIN=CB; TISSUE=Bursa;
Caldwell R.B., Kierzek A.
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NON_TER
SEQUENCE 36
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N5 CHICK
Q5ZKN5 CHICK
Q5ZKN5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN-CB; TIŜSUE-Bursa;
Caldwell R.B., Klerzek A.M., Arakawa H., Bezzubov Y Fiedler P., Kuterz S., Blagodatski A., Kostovska D. Plachy J., Carninci P., Hayashizaki Y., Buerstedde "Full-length cDNAs from chicken bursal lymphocytes genefunction analysis";
Genome Biol. 6:R6-R6(2005).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-MAR-2005, integrated into t
15-MAR-2005, sequence version
07-FEB-2006, entry version 3.
Hypothetical protein.
ORFNames=RCJMB04_33c19;
                                                                                                                                                                                                                       23-NOV-2004, integrated into UniProtKB/TrEMBL. 23-NOV-2004, sequence version 1. 07-FBB-2006, entry version 6. Hypothetical protein.
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Q5F360;
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Gallus gallus (Chicken)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hypothetical SEQUENCE 4:
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Archosauria; Aves; Neognathae;
                                                                                                                                              Archosauria;
                                                                                                                                                                  Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Distributed under the Creative Commons
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Attribution-NoDerivs License
          Kostovska
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KQR6 PLACH

QAXQR6 PLACH

PRELIMINARY; P.

QAXQR6;

05-JUL-2005, integrated into Un

05-JUL-2005, sequence version 1

21-FEB-2006, entry version 7.

Hypothetical protein.

ORFNames=PC000206.04.0;
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QBQHJ7;
01-JUN-2002, integrated into U
01-JUN-2002, sequence version
07-FEB-2006, entry version 8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Flaciny J., Carninci P., Hayashizaki Y., "Full-length cDNAs from chicken bursal ] genefunction analysis.";
Genome Biol. 6:R6-R6(2005).
                                                                                                                                                                                                                                                                                                                                                            MEDLINE=22021057; PubMed=11984880; DOI=10.1002/dvdy.10090;
Jun L., Balboni A.L., Laitman J.T., Bergemann A.D.;
"Isolation of DNTNP, which encodes a potential nuclear pro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Dorsal neural-tube Name=DNTNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hypothetical SEQUENCE 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Copyrighted by the UniProt Consortium, Distributed under the Creative Commons
          Plasmodium chabaudi.
Eukaryota; Alveolata;
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Distributed
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Commons Attribution-NoDerivs License
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                                                                                                                                                                STRAIN=CO-92 / Blovar Orientalis;

MEDLINE=21470413; PubMed=11586360; DOI=10.1038/35097083;

Parkhill J., Wren B.W., Thomson N.R., Titball R.W., Holden M.T.G.,

Prentice M.B., Sebaihia M., James K.D., Churcher C.M., Mungall K.L.,

Baker S., Basham D., Bentley S.D., Brooks K., Cerdeno-Tarraga A.-M.,

Chillingworth T., Cronin A., Daviss R.M., Davis P., Dougan G.,

Feltwell T., Hamlin N., Holroyd S., Jagels K., Karlyshev A.V.,

Leather S., Moule S., Oyston P.C.F., Quail M.A., Rutherford K.M.,

Simmonds M., Skelton J., Stevens K., Whitehead S., Barrell B.G.;

"Genome sequence of Yersinia pestis, the causative agent of plague.",

Nature 413:523-527(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                             MAO1 YERPE STANDARD; PRT; 565 AA.

(082G09; 074VB1; 07CHQ5;

06-DEC-2005, integrated into UniProtKB/Swiss-Prot
01-MAR-2002, sequence version 1.

07-MAR-2006, entry version 27.

NAD-dependent malic enzyme (EC 1.1.1.38) (NAD-ME)
Name=sfck; OrderedLocusNames=YPO1511, y2658, YP14
             STRAIN=KIM5 / Biovar Mediaevalis;
MEDLINE=22137863; PubMed=12142430;
DOI=10.1128/JB.184.16.4661.4611.2002;
Deng W., Burland V., Plunkett G. III, Boutin A., Mayhew
Perna N.T., Rose D.J., Mau B., Zhou S., Schwartz D.C.,
Fetherston J.D., Lindler L.E., Brubaker R.R., Plano G.V.
Straley S.C., McDonough K.A., Nilles M.L., Matson J.S.,
Perry R.D.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUCLEOTIDE SEQUENCE.

PubMed=15637271; DOI=10.1126/science.1103717;

Hall N., Karras M., Raine J.D., Carlton J.M., Kooij T.W.A.,

Hall N., Karras M., Raine J.D., Carlton J.M., Christophides G.K.,

James K., Rutherford K., Harris B., Harris D., Churcher C.M.,

James K., Rutherford K., Harris B., Harris D., Churcher C.M.,

Quail M.A., Ormond D., Doggett J., Trueman H.E., Mendoza J.,

Bidwell S.L., Rajandream M.A., Carucci D.J., Yates J.R. III,

Kafatos F.C., Janse C.J., Barrell B.G., Turner C.M.R., Waters A.P.,
                                                                                                                                                  NUCLEOTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           transcriptomic, and pro
Science 307:82-86(2005)
                                                                                                                                                                                                                                                                                                                                                                                                                    Enterobacteriaceae; Yersinia.
                                                                                                                                                                                                                                                                                                                                                                                                                                     Bacteria;
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                                                                                                                                                                                                                                                                                                                                                                   NUCLEOTIDE SEQUENCE [LARGE SCALE
                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=632;
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GO:0005615; C:extracellular space; IEA.
GO:0005125; F:cytokine activity; IEA.
erPro; IPR012351; Cytokine 4 hix.
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GenomeReviews; ABO09952 GR; YP1401.
GenomeReviews; AL590842 GR; YP01511.
BioCyc; YPES187410:Y2658-MONMOMER; --
BioCyc; YPES29193:YP1401-MONOMER; --
BioCyc; YPES32:YP01511-MONOMER; --
HAMAP; MF_01619; -; 1.
InterPro; IPR012301; Malic_NAD_bd.
InterPro; IPR012302; Malic_NAD_bd.
InterPro; IPR012302; Malic_Oxred.
PANTHER; PTHR10893; Malic_Oxred; 1.
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NAD-dependent malic
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                                                                                                                                                                                 SEQUENCE
                                                                                                                                                                                                                                                                                                                                   Pfam; PF00390; malic; 1. Pfam; PF03949; Malic M; PIRSF; PIRSF000106; ME;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Copyrighted by the UniProt Consortium, Distributed under the Creative Commons
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        J. Bacteriol. 184:4601-4611(2002)
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AE017132; AAS61642.1;
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Proc. Natl. Acad. Sci. U.S.A. 101:13826-13831(2004).

-!- COFACTOR: Divalent cations, magnesium or manganese (By
                                                                                                                        Q6CS90;
20-DEC-2005, integrated into Unif
16-AUG-2004, sequence version 1.
07-FEB-2006, entry version 14.
Diphthamide biosynthesis protein
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InterPro; IPR012301; Malic N.
InterPro; IPR01302; Malic_NAD_bd.
InterPro; IPR001891; Malic_oxred.
PANTHER; PTHR.10893; Malic_oxred; 1.
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                                                         Name=DPH2; OrderedLocusNames=KLLA0D02948g; Kluyveromyces lactis (Yeast).
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   Saccharomycetales;
                              Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
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-!- SIMILARITY: Belongs to the malic enzymes family.
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PubMed=15358858; DOI=10.1073/pnas.0404012101;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PF00390; malic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BX936398;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 DYKDDDDK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           DFKNDDDK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   565
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                             STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           73
   Saccharomycetaceae;
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565 NAD-dependent malic enzyme.
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NAD (By similarity).
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Pred. No. 8.9e+02;
2; Mismatches 0;
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F57C1183EEC4AA3D CRC64;
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Proton acceptor (By similarity)
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                                                                                                                                                                                                                                                                                             PRT;
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   Kluyveromyces.
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RESULT
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Lafontaine I., de Montigny J., Marck C., Neuveglise C., Talla E.,
Goffard N., Frangell L., Alpide M., Anthouard V., Babour A., Barbe V.
Barnay S., Blanchin S., Beckerich J.-M., Beyne E., Bleykasten C.,
A. Boisrame A., Boyer J., Cattolico L., Confanioleri F., de Daruvar A.
Despons L., Fabre E., Fairhead C., Ferry-Dumazet H., Groppi A.,
A. Hantraye F., Hennequin C., Jauniaux N., Joyet P., Kachouri R.,
A. Kerrest A., Koszul R., Lemaire M., Lesur I., Ma L., Muller H.,
A. Kerrest A., Koszul R., Lemaire S., Ozier-Kalogeropoulos O.,
A. Nicaud J.-M., Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,
A. Nicaud J.-M., Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,
A. Pellenz S., Potier S., Richard G.-F., Straub M.-L., Suleau A.,
A. Pellenz S., Potier S., Richard G.-F., Straub M.-L., Suleau A.,
A. Swennen D., Tekaia F., Wesolowski-Louvel M., Westhof E., Wirth B.,
Zeniou-Meyer M., Zivanovic Y., Bolotin-Fukuhara M., Thierry A.,
B. Bouchier C., Caudron B., Scarpelli C., Gaillardin C., Weissenbach C.,
A. Wincker D. C., Caudron B., Scarpelli C., Gaillardin C., Weissenbach C.,
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Best Local S
Matches 6
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:B4 DICDI
  Q55CB4 DICDI
  Q55CB4;
  24-MAY-2005;
PubMed=15875012; DOI=10.1038/nature03481; Bichinger L., Pachebat J.A., Gloeckner G., Rajandream M.A., Sucgang R., Berriman M., Song J., Olsen R., Szafranski K., Xu Q., Sucgang B., Kummerfeld S., Madera M., Konfortov B.A., Rivero F., Bankier A.T., Lehmann R., Hamlin N., Davies R., Gaudet P., Fey P., Pilcher K., Chen G., Saunders D., Sodergren E.J., Davis P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; CR382124; CAH00295.1; -; Genomic InterPro; IPR002728; Diphthamide_syn. InterPro; IPR010014; DPH2. PTHR10762; Diphthamide_syn; PANTHER, PTHR10762; Diphthamide_syn; 1. TIGR07848; TIGR00272; DPH2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC STRAIN=CBS 2359 / IFO 1267 / NRRL Y-1140 PubMed=15229592; DOI=10.1038/nature02579;
                                                                                                                                                                                                                      07-FEB-2006, entry v
Hypothetical protein
ORFNames=DDB0190829;
                                                                                                                                                           Eukaryota; Mycetozoa;
NCBI_TaxID=44689;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Copyrighted by the UniProt Consortium, see http://www.uniprot.org/tonstributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bouchier C., Caudron B., Scarpelli Wincker P., Souciet J.-L.;
                                                                                                             STRAIN=AX4
                                                                                                                                                                                                   Dictyostelium discoideum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CHAIN 1
                                                                                                                                NUCLEOTIDE
                                                                                                                                                                                                                                                      24-MAY-2005, sequence version 07-FEB-2006, entry version 4.
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nes 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SUBCELLULAR LOCATION: Cytoplasm (By similarity). SIMILARITY: Belongs to the DPH2 family.
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                                                                                                                                                                                                                                                                                                                                                                                                                           553
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85.7%;
                                                                                                                                                                                  pum (Slime mold).
Dictyosteliida;
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Pred. No. 9.3e+02;
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                                                                                                                                                                                                                                                                                                UniProtKB/TrEMBL.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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/ WM37;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 587;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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Barbe
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RESULT 40
Q6BYD0 DEBHA
ID Q6BYD0;
DT 16-AUG-2
DT 16-AUG-2
DT 07-FEB-2
CO Debaryom
DE hansenii
GN 07deredi
OS Debaryom
DC Saccharo
OC Saccharo
OC NCBI_Tax
RN (1)
RC STRAIN-A
RX PUBMed-1
RC STRAIN-A
RX PUBMed-1
RA Dujon B.
RA Lafontai
RA Goffard
RA Dujon B.
RA Lafontai
RA Goffard
RA Barnay S
RA Boisrame
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RA Hantraye
RA Kerrest
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              NUCLEOTIDE SEQUENCE (LARGE SCALE GENOMIC DNA).

RETRAIN=ATCC 36239 / CBS 767;

RETRAIN=ATCC 3
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Best Local
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16-AUG-2004,
07-FEB-2006,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kerhornou A., Nie X., Hall N., Anjard C., Hemphill L., Bason N., Farbrother P., Desany B., Just E., Morio T., Rost R., Churcher C.M., Cooper J., Haydock S., van Driessche N., Cronin A., Goodhead I., Muzny D.M., Mourier T., Pain A., Lu M., Harper D., Lindsay R., Hauser H., James K.D., Quiles M., Madan Babu M., Saito T., Buchrieser C., Wardroper A., Felder M., Thangavelu M., Johnson D., Knights A., Loulseged H., Hernandez J., Rabbinowitsch E., Steffen D., Quail M.A., Urushihara H., Hernandez J., Rabbinowitsch E., Steffen D., Sanders M., Ma J., Kohara Y., Sharp S., Simmonds M.N., Spiegler S., Tivey A., Sugano S., White B., Walker D., Woodward J.R., Winckler T., Tanaka Y., Shaulsky G., Schleicher M., Weinstock G.M., Rosenthal A., Cox E.C., Chisholm R.L., Gibbs R.A., Loomis W.F., Platzer M., Kay R.R., Williams J.G., Dear P.H., Noegel A.A., Barrell B.G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Debaryomyces hansenii (Yeasī) (Yorulaspora hansenii).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Debaryomyces.
NCBI_TaxID=4959;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pfam; PF01485; IBR; 1.
Pfam; PF05773; RWD; 1.
SMART; SM00647; IBR; 1.
SMART; SM00591; RWD; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              InterPro;
InterPro;
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"The c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OrderedLocusNames=DEHA0A10868g;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            QEBADO_DEBHW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hypothetical SEQUENCE 58
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Distributed
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XA Machida M., Asai K., Sano M., Tanaka T., Kumagai T., Terai G.,
XA Machida M., Asai K., Sano M., Tanaka T., Xumagai T., Gomi K.,
XA Kusumoto K., Arima T., Akita O., Kashiwagi Y., Abe K., Gomi K.,
XA Horiuchi H., Kitamoto K., Kobayashi T., Takeuchi M., Denning D.W.,
XA Galagan J.E., Nierman W.C., Yu J., Archer D.B., Bennett J.W.,
XA Galagan J.E., Nierman W.C., Yu J., Archer D.B., Bennett J.W.,
XA Hosoyama A., Ichinomiya M., Igarashi R., Iwashita K., Juvvadi P.R.,
XA Hosoyama A., Ichinomiya M., Igarashi R., Iwashita K., Juvvadi P.R.,
XA Hosoyama A., Ichinomiya M., Igarashi R., Iwashita K., Dulsen I.,
XA Kato M., Kato Y., Kin T., Kokubun A., Maeda H., Maeyama N.,
XA Maruyama J., Nagasaki H., Nakajima T., Oda K., Okada K., Paulsen I.,
XA Kamorto K., Sawano T., Takahashi M., Takase K., Terabayashi Y.,
XA Sakamoto K., Sawano T., Takahashi M., Takase K., Terabayashi Y.,
XA Komori T., Koyama Y., Minetoki T., Suharnan S., Tanaka A., Isono K.,
XA Kuhara S., Ogasawara N., Kikuchi H.;
Yenome sequencing and analysis of Aspergillus oryzae.";
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Nature 438:1157-1161(2005).
                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN=RIB
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Eukaryota; Fungi; Ascomycot;
Eurotiales; Trichocomaceae;
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ORFNames=A0090023000761;
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A Kusumoto K., Arima T., Akita O., Kashiwagi Y., Abe K., Gomi K.,

A Horiuchi H., Kitamoto K., Kobayashi T., Takeuchi M., Denning D.W.,

A Horiuchi H., Kitamoto K., Kobayashi T., Takeuchi M., Denning D.W.,

A Galagan J.E., Nierman W.C., Yu J., Archer D.B., Bennett J.W.,

A Hosoyama J., Cleveland T.E., Fedorova N.D., Gotoh O., Horikawa H.,

A Hosoyama A., Ichinomiya M., Igarashi R., Iwashita K., Juvvadi P.R.,

A Hosoyama A., Ichinomiya M., Marashi R., Maeda H., Maeyama N.,

A Kato W., Kato Y., Kin T., Kokubun A., Maeda H., Maeyama N.,

A Maruyama J., Nagasaki H., Nakajima T., Oda K., Okada K., Paulsen I.,

A Maruyama J., Nagasaki H., Nakajima T., Oda K., Okada K., Paulsen I.,

Sakamoto K., Sawano T., Takahashi M., Takase K., Terabayashi Y.,

Sakamoto K., Sawano T., Takahashi M., Takase K., Terabayashi Y.,

Sakamoto T., Koyama Y., Minetoki T., Suharnan S., Tanaka A., Isono K.,

A Komori T., Koyama Y., Minetoki H.,

"Genome sequencing and analysis of Aspergillus oryzae.";

"Genome sequencing and analysis of Aspergillus oryzae.";
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Bacteria; Proteobacteria; Alphaproteobacteria;
Rhodobacteraceae; Silicibacter.
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Machida M., Asai K., Sano M., Tanaka T.,
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                                                                           "Sequencing TM1040.";
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ORFNames=RoseDRAFT_18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DYSDDDDDQ
                                                                                           A., Lucas S., Lapidus A., Barry K., ., Israni S., Pitluck S., Richardson ing of the draft genome and assembly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           рукрарарк в
                                                 (JUL-2005)
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F:kinase activity; IEA.
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1.3e+03;
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                                                                                           P.;
of Sili
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Isono K.,
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US DOE Joint Genome Institute (JGI-PGF);
US DOE Joint Genome Institute (JGI-PGF);
Copeland A., Lucas S., Lapidus A., Barry K., Detter C.,
Copeland A., Lucas S., Pitluck S., Richardson P.;
Hammon N., Israni S., Pitluck S., Richardson P.;
Submitted (JUN-2005) to the EMBL/GenBank/DDBJ databases
-i- CAUTION: The sequence shown here is derived from an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US DOB Joint Genome Insti
Larimer F., Land M.;
"Annotation of the draft
Submitted (JUL-2005) to t
                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=22549451; PubMed=12663533; Astrom S.U., Cline T.W., Rine J.; "The Drosophila melanogaster sir2(+) Gene Is Nonessential Minor Effects on Position-Effect Variegation."; Genetics 163:931-937(2003).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-MAY-1999, integrated into
01-MAY-1999, sequence versior
07-FEB-2006, entry version 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hypothetical protein; Signal SIGNAL 1 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Copyrighted by the UniProt Consortium, see http://www.uniprot.org/tubistributed under the Creative Commons Attribution-NoDerivs License
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Neoptera; Endopterygota; Dip
Ephydroidea; Drosophilidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Drosophila 
Eukaryota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SIR2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBL; AAFG02000004; EAN57069.1; -;
                                                                                                                                                                                                                                                                                                                                               Copyrighted by the UniProt Consortium, see http://www.uniprot.Distributed under the Creative Commons Attribution-NoDerivs Li
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Name=Sir2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUCLEOTIDE
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                                                                                      SEQUENCE
                                                                                                         Pfam; PF02146;
PROSITE; PS5030
                                                                                                                                                    InterPro; IPR003000; SIR2.
PANTHER; PTHR11085; SIR2; 1.
                                                                                                                                                                                                                                                                  FlyBase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUCLECTIDE SEQUENCE.
                                                                                                                                                                                             Base; FBgn0024291; Sir2.
GO:0005737; C:cytoplasm; IDA
GO:0005634; C:nucleus; IDA.
GO:0017136; F:NAD-dependent
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                                                                                                                                                                                                                                                                                      P53686;
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Similarity 6; Conserv
                                                                                                         02146; SIR2;
PS50305; SI
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melanogaster (Fruit fly).
Metazoa; Arthropoda; Hexapoda;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
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                                                                                                                                                                                                                                                                                                           AAC79684.1;
                                                                                    SIRTUIN; 1.
; 92024 MW;
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ae; Drosophila
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  Score 38; DB 2;
Pred. No. 1.3e+03;
1; Mismatches 1
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                                                                                                                                                                                                                                                                                                               mRNA.
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                                                                                         5EF09F1A46235E7F CRC64;
                                                                                                                                                                                                  histone deacetylase activity; IDA.
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                                             Length 823;
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RA Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D.,
RA Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Goalle R.F.,
RA George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N.,
RA George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N.,
RA Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
RA Burtin G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
RA Burtin G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
RA Burtin G.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D.,
RA Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M.,
RA Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M.,
RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,
RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier R.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Gebablos B., Delcher A., Deng Z., Mays A.D., Dew I., Dietz S.M.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Godson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,
RA Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M.,
RA Harris N.L., Harvey D.A., Heiman T.J., Hermandez J.R., Houck J.,
RA Harris N.L., Harvey D.A., Heiman T.J., Hermandez J.R., Houck J.,
RA Harris N.L., Mattei B., McIncosh T.C., McLeod M.-H., Ingewan C.,
RA Hasko P., Lei Y., Levitsky A.A., Li J.H., Li Z., Liang Y., Lin X.,
RA McIston D.R., Nelson K.A., Saunders R.D. C., Scheeler F., Shen H.,
RA Merkulov G., Milshina N.V., Mobarry C., Morris J., Woshrefi A.,
RA Melson D.R., Nelson K., Nixon K., Nuskern D.R., Pacieb J.M.,
RA Melson D.R., Nelson K., Nuskern D.R., Pacieb J.M.,
RA Meng E.C., Stapleton M., Strong R., Sun E.,
RA Svirskas R., Tector C., Turner R., Venter T., Wang A.H., Wang X.,
RA Spier B., Spradling A.C., Stapleton M., Strong R., Sun E.,
RA Gibbs R.A., Wers E.W., Rubin G.M., Zhang G., Zhao Q., Zheng L.,
RA Gibbs R.A., Wers E.W., Rubin G.M., Zhou X., Zhao Q., Zheng L.,
Ra Gibbs R.A., Mers E. S., Scholl R., Scholl H.O.,
Ra G
                                                                                                                                                                                                                                                                                             Zheng X.H., Keh K
Gibbs R.A., M
NUCLEOTIDE SEQUENCE.
Nucleotide S., Adams M., Champe M., Dugan S.P., Frise E., Hodgson A., Patel S., Adams M., Champe M., Dugan S.P., Muzny D.M., Nelson C.R., George R.A., Hoskins R.A., Laverty T., Muzny D.M., Nelson E.J., Pacleb J.M., Park S., Pfeiffer B.D., Richards S., Sodergren E.J., Pacleb J.M., Park S., Pfeiffer B.D., Richards S., Sodergren E.J., Sylrskas R., Tabor P.E., Wan K., Stapleton M., Sutton G.G., Venter (Weinstock G., Scherer S.E., Myers E.W., Gibbs R.A., Rubin G.M., "Finishing a whole-genome shotgun: release 3 of the Drosophila "Finishing a whole-genome shotgun: release 3 of the Drosophila
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.
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07-FEB-2006,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PubMed=10731132; DOI=10.1126/science.287.5461.2185;
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PANTHER; PTHR11085; SIR2; 1.
Pfam; PF02146; SIR2; 1.
PROSITE; PS50305; SIRTUIN; 1.
CROTTENCE 823 AA; 91837 MW;
                               Q4Y0Z6 PLACH
Q4Y0Z6;
05-JUL-2005,
05-JUL-2005,
07-FEB-2006,
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           Asparagine-rich antigen Pfa55-14, ORFNames=PC000038.02.0;
                                                                                                                                                                                                                                                                                                                                                               EMBL; AE003639; AAFE
                                                                                                                                                                                                                                                                                               888
                                                                                                                                                                                                                                                                                                                                                                                                     Copyrighted by the UniProt Consortium, see http://www.uniprot.org/teDistributed under the Creative Commons Attribution-NoDerivs License
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Celniker S., Carlson J.,
Hoskins R., Stapleton M.,
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systematic review."
                                                                                                                                                                                                                                                                                                                                             FlyBase;
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MEDLINE=22426069; PubMed=12537572;
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Submitted (MAR-2000) to t
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Genome Biol. 3:RESEARCH0084.1-RESEARCH0084.20(2002).
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MEDLINE=22426070; PubMed=12537573;
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PLACH
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GO:0005737; C:cytoplasm; IDA.
GO:00005734; C:nucleus; IDA.
GO:0017136; F:NAD-dependent histone
GO:0005515; F:protein binding; IPI.
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D.A., Lewis S.E.,
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Svirskas R.,
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Matches 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].

STRAIN=CLIB 122 / E 150;

PubMed=15229502; DOI=10.1038/nature02579;

Pubmed=15229502; DOI=10.1038/nature02579;

Dujon B., Sherman D., Fischer G., Durrens P., Casaregola Dujon B., Sherman D., Fischer G., Menveglise C., Tafontaine I., de Montigny J., March C., Neuveglise C., The Control of the Control of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OrderedLocusNames=YALIOB00528g;
Yarrowia lipolytica (Candida lipolytica).
Eukaryota; Fungi; Ascomycota; Saccharomycotina;
Saccharomycetales; Dipodascaceae; Yarrowia.
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21-FEB-2006, entry version 19.
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PubMed=15637271; DO
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NCBI_TaxID=5825;
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Nature 430:35-44(2004).
                                                                                                                                                                                                                                                    Wincker P.,
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                                                                                                                   SUBCELLULAR LOCATION: Cytoplasm
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P., Souciet J.-L.;
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M., Raine J.D., Carlton J.M.,
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elli C., Gaillardin C., Weissenba
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RA AFACRICAL H.M., January C., Calvo S.E., Camarata J., Chang J., Rh Boukhgalter B., Butler J., Cook A., Cooke P., Corum B., DeArellano K., Rh Choepel Y., Collymore A., Cook A., Cooke P., Corum B., DeArellano K., Rh Choepel Y., Collymore A., Cook P., Corum B., DeArellano K., Rh Choepel Y., Collymore A., Cook P., Corum B., DeArellano K., Rh Choepel Y., Collymore A., Cook P., Corum B., DeArellano K., Rh Choepel Y., Dodge S., Dooley K., Dorris L., Elkins T., Engels R., Rh Checkson J., Faro S., Ferreira P., FitzGerald M., Gage D., Rh Carthy M., Camat M., Gage D., Parker N., Rh Hagopian D., Hagos B., Hall J., Horton L., Hulme W., Rh Hafez N., Hagopian D., Hagos B., Hall J., Horton L., Hulme W., Rh Hafez N., Hagopian D., Manders T., Levine R., Lindblad-Toh K., Liu G., Rh L.-J., Mabbitt R., MacLean C., Macdonald P., Major J., Rh Karatas A., Kells C., Landers T., Levine R., Lindblad-Toh K., Liu G., Rh Manning J., Matthews C., Mauceli E., McCarthy M., Meldrim J., Rh Manning J., Matthews C., Mauceli E., McCarthy M., Meldrim J., Rh Manning J., Matthews C., Mauceli E., McCarthy M., Meldrim J., Rh Manning J., Milova T., Mlenga V., Murphy T., Naylor J., Nguyen C., Rh Micol R., Nielsen C.B., Norbu C., O'Connor T., O'Donnell P., Rh O'Neil D., Oliver J., Peterson K., Phunkhang P., Pierre N., Rh O'Neil D., Oliver J., Peterson K., Phunkhang P., Pierre N., Rh Rogov P., Roman J., Schauer S., Schupback R., Seaman S., Severy P., Rh Smirnov S., Smith C., Spencer B., Stange-Thomann N., Stojanovic N., Rh Stubbs M., Talamas J., Tesfaye S., Theodore J., Topham K., Travers M., Vassiliev H., Venkataraman V.S., Viel R., Vo A., Wang S., Wilson B., Whith M., Wang S., Walson B., Walson B., W
RESULT PROCESS OF PROC
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Gibberella zeae (Pusarium graminearum).
Bukaryota; Pungi; Ascomycota; Pezizomycotina; Sordariomycetes;
Bukaryota; Pungi; Ascomycota; Nectriaceae; Gibberella.
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GO; GO:0005488; F: binding; IEA.
InterPro; IPR01199; TPR-like_helical.
InterPro; IPR011440; TPR 1.
InterPro; IPR013105; TPR 2.
InterPro; IPR013105; TPR 2.
InterPro; IPR013105; TPR 2.
InterPro; IPR013026; TPR_region.
Pfam; PF00515; TPR 1; 4.
SMART; SM00028; TPR; 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Birren B.W., Nusbaum C., Abouelleil A., Allen N., Arachchi H.M., Barna N., Bastien V., Bloom T., Bookhgalter B., Butler J., Calvo S.E., Camarata J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-AUG-2005, integrated into 16-AUG-2005, sequence version 07-FEB-2006, entry version 4.
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Q416J9;
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PROSITE; PS50293; TPR; TSCION; 3
Complete proteome; Repeat; TPR
SEQUENCE 926 AA; 105464 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hypothetical protein.
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                                                                                                                                            Copyrighted by the UniProt Consortium, Distributed under the Creative Commons
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                                                                                                                                                                                                                                                                                     CAUTION: The sequence shown here is derived EMBL/GenBank/DDBJ whole genome shotgun (WGS)
                                                                                                                                                                                                                                                   preliminary data.
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85.7%;
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0894AED20AD98CD3 CRC64;
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EMBL; AACM01000301; Complete proteome; H SEQUENCE 1018 AA;

Hypothetical ; 114167 MW;

EAA76618.1; -; Genomic

DNA

Attribution-NoDerivs License

protein

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RESULT 49
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A Johnston M., Hillier L.W., Riles L., Albermann K., Andre B.,
A Johnston M., Hillier L.W., Riles L., Delius H., Dubois E.,
A Ansorge W., Benes V., Brueckner M., Delius H., Dubois E.,
A Duesterhoeft A., Entlan K.-D., Floeth M., Goffeau A., Hebling U.,
A Heumann K., Heuss-Neitzel D., Hilbert H., Hilger F., Kleine K.,
A Koetter P., Louis E.J., Messenguy F., Mews H.-W., Miosga T.,
A Koetter D., Mueller Auer S., Nentwich U., Obermaier B., Piravandi I.
A Moestl D., Mueller S., Purnelle B., Rechmann S., Rieger M.,
A Pohl T.M., Portetelle D., Purnelle B., Rechmann S., Rieger M.,
A Rinke M., Rose M., Scharfe M., Scherens B., Scholler P., Schwager A.,
A Rinke M., Rose M., Scharfe M., Vorestarazu L.A., Vandenbol M.,
A Verhasselt P., Vierendeels F., Voet M., Volckaert G., Voss H.,
A Wambutt R., Wedler E., Wedler H., Zimmermann F.K., Zollner A.,
A Hani J., Höheisel J.D.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                  MEDLINE=22923965; PubMed=14562106; DOI=10.1038/nature02046; Ghaemmaghami S., Huh W.-K., Bower K., Howson R.W., Belle A., Dephoure N., O'Shea E.K., Weissman J.S.; "Global annalysis of protein expression in yeast."; Nature 425:737-741(2003).
-I- FUNCTION: Controls the pre-rRNA processing machinery in conjunction with FHL1. Could convert FHL1 from a repress
                                        PIR; S55352; S55352.
                                                                                            Copyrighted
Distributed
                                                                                                                                                                                                                                                                                                     LEVEL OF PROTEIN EXPRESSION.
                                                                                                                                                                                                                                                                                                                             "The nucleotide sequence of Saccharomyces Nature 387:87-90(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN=S288c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Saccharomyces cerevisiae.",
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRAIN=ATCC 28383 / FL100;
MEDLINE=95304839; PubMed=7785326;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Saccharomyces cerevisiae (Baker's yeast).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Sacci
Saccharomycetales; Saccharomycetaceae; Saccharomyces
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-MAR-2006,
Protein IFH1
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01-FEB-1995,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUCLEOTIDE SEQUENCE
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P39520;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUCLEOTIDE SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=4932;
                                                                                                                                               activator.
INTERACTION:
Q08673:SRL1; NbExp=1;
SUBCELLULAR LOCATION:
                                                                                                                                    MISCELLANEOUS: Present with 1430 molecules/cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           el I., Thuriaux P.;
IFH1 gene product interacts
                                                  Z29488; CAA82624.1;
U19027; AAB67412.1;
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                         P39520;
YLR223C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ασσσαγα
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                                                                                            by the Uni
              142285;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         integrated into UniProtKB/Swiss-Prot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AB972
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Saccharomyces cerevisiae
                                                                                                          UniProt
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FL100;
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80
                                                                                            iProt Consortium, see http://www.uniprot.org/t.
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                                                                                                                                                  Nucleus
                                                                                                                                                               IntAct=EBI-9054,
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                                                Genomic_DNA.
Genomic_DNA.
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RA Cuomo C., David R., Dasvoe T., Degray S., Dodge S., Dooley K.,
RA Dorje P., Dorjee K., Dorris L., Duffey N., Dupes A., Elkins T.,
RA Engels R., Erickson J., Farina A., Faro S., Ferreira P., Fischer H.,
RA Enizgerald M., Foley K., Gage D., Galagan J.E., Gearin G., Gnerre S.,
RA Gnirke A., Goyette A., Graham J., Grandbois E., Gyaltsen K., Hafez N.,
RA Hagopian D., Hagos H., Hall J., Hatcher B., Heller A., Higgins H.,
RA Honan T., Horn A., Houde N., Hughes L., Hulme W., Hugby E., Iliev I.,
RA Jaffe D., Jones C., Kamal M., Kamat A., Kamvysselis M., Karlsson E.,
RA Kells C., Kieu A., Kisner P., Kodira C., Kulbokas E., Labutti K.,
RA Lama D., Landers T., Leyer J., Levine S., Lewis T.,
RA Lindblad-Toh K., Liu X., Lokyitsang T., Lokyitsang Y., Lucien O.,
RA Manning J., Marabella R., Maru K., Matchews C., Mauceli E.,
RA McCarthy M., McDonough S., McGhee T., Meldrim J., Meneus L.,
RA McCarthy M., McDonough S., McGhee T., Meldrim J., Meneus L.,
RA Mozes J., Mulrain L., Munson G., Naylor J., Newes C., Nauceli E.,
RA Mozes J., Mulrain L., Munson G., Nallelsen T., Mlenga V., Moru K.,
RA Mozes J., Mulrain L., Munson G., Nalles R., Nielsen C.B., Nizzari M., Norbu C.,
RA Norbu N., O'Donnell P., Okoawo O., O'Leary S., Contosho B.,
RA O'Neill K., Osman S., Parker S., Perrin D., Phunkhang P., Piqani B.,
RA Purcell S., Rachupka T., Ramasamy U., Rameau R., Ray V., Raymond C.,
RA Retta R., Richardson S., Rise C., Rodriguez J., Rogers J., Rogov P.,
RA Sheridan J., Sherpa N., Shi J., Smirnov S., Smith C., Sougnez C.,
RA Spencer B., Stalker J., Stange-Thomann N., Stavropoulos S.,
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Best Local
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19-JUL-2005,
07-FEB-2006,
Hypothetical
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SEQUENCE
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LinkHub; P39520; -
G0; G0:0005730; C:nucleolus; IDA.
G0; G0:0005515; F:protein binding; IPI.
G0; G0:0005515; F:transcription factor activity;
G0; G0:0003700; F:transcription factor activity;
G0; G0:0006348; P:chromatin silencing at
G0; G0:0006364; P:rRNA processing; IGI.
G0; G0:0006363; P:transcription from RNA polymer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Calvo S.E.,
Citroen M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Birren B.W., Nusbaum C., Abebe A., Abouerrerr B.W., Nusbaum C., Abebe A., Abouerrerr B.W., Anderson S., Ait-Zahra M., Allen N., Allen T., An P., Anderson M., Anderson S., Arachchi H.M., Armbruster J., Bachantsang P., Baldwin J., Barry A., Arachchi H.M., Armbruster J., Bloom T., Blye J., Boguelavskiy L., Pavul T., Blitshsteyn B., Bloom T., Blye J., Brutler J., Calixte N.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAIN=521;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID≃237631;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ustilāginomycetidae;
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Q4P6M8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUCLEOTIDE SEQUENCE [LARGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Fungi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Transcription
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GO:0005515; F:protein binding; IPI.
GO:0003700; F:transcription factor activity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              S000004213;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Collymore A., Considine T., Cook A., Cooke
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           , Boukhgalter B., Brunache A., Butler J., Calixte Camarata J., Campo K., Chang J., Cheshatsang Y.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  122
1085
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     integrated into UniProtKB/TrEMBL.
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ae; Ustilaginales; Ustilaginaceae;
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1, 122492
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85.7%;
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/FTId=PRO_
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Asp/Glu-rich (highly
W; BE1C7DEF06213FE0
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CRC64;
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Iliev I.,
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Matches 6
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GO; GO:000534; C:nucleus; IEA.
GO; GO:0046872; F:metal ion binding; IEA.
GO; GO:00048720; F:metal ion binding; IEA.
GO; GO:0003700; F:transcription factor activity; IEA.
GO; GO:0008270; F:zinc ion binding; IEA.
GO; GO:0008350; P:regulation of transcription, DNA-dependent; IEA.
GO; GO:0006350; P:transcription; IEA.
InterPro; IFRO01138; Fungi_Trscrp_N.
                                                                                                                                     Dewhirst F.E., Fraser C.M.;
"Complete genome sequence of the oral pathogenic bacterium Porphyromonas gingivalis strain W83.";
J. Bacteriol. 185:5591-5601(2003).
-!- CATALYTIC ACTIVITY: Deoxynucleoside triphosphate + DNA(n)
                                                                                                                                                                                                                                                                                                                      DOI=10.1128/JB.185.18.5591-5601.2003;
Nelson K.E., Fleischmann R.D., Deboy R.T., Paulsen I.T., Fouts D.E.,
Eisen J.A., Daugherty S.C., Dodson R.J., Durkin A.S., Gwinn M.L.,
Haft D.H., Kolonay J.F., Nelson W.C., Mason T.M., Tallon L., Gray J.
Granger D., Tettelin H., Dong H., Galvin J.L., Duncan M.J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PROSITE; PS00463; ZN2_CY6_FUNGAL_1; UNKNOWN_1.
PROSITE; PS50048; ZN2_CY6_FUNGAL_2; 1.
DNA-binding; Hypothetical protein; Metal-binding; Nuclear Transcription; Transcription regulation; Zinc.
SEQUENCE 1212 AA; 131152 MW; 7182B4E422BFF507 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "The genome sequence of Ustilago maydis.";
Submitted (FEB-2004) to the EMBL/GenBank/DDBJ da
-!- CAUTION: The sequence shown here is derived
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Stetson K., Stone C., Stone S., Stubbs M., Talamas J., Tchuinga P., Tenzing P., Tesfaye S., Theodore J., Thoulutsang Y., Topham K., Towey S., Tsamla T., Tsomo N., Vallee D., Vassiliev H., Venkataraman V.S., Vinson J., Vo A., Wade C., Wang S., Wangchuk T., Wangdi T., Whittaker C., Wilkinson J., Wu Y., Wyman D., Yadav S., Yang S., Yang X., Yeager S., Yee E., Young G., Zainoun J., Zembeck L., Zimmer A., Zody M., Lander E.S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Porphyromonas gingivalis (Bacteroides gingivalis)
Bacteria; Bacteroidetes; Bacteroidetes (class); E
Porphyromonadaceae; Porphyromonas.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-DEC-2003, integrated into UniProtKB/TrEMBL
15-DEC-2003, sequence version 1.
21-FEB-2006, entry version 22.
DNA polymerase III, alpha subunit.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pfam; PF04082; Fungal_trans
Pfam; PF00172; Zn_clus; 1.
SMART; SM00066; GAL4; 1.
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   Copyrighted
                                                                      diphosphate + DNA(n+1).
-!- SUBCELLULAR LOCATION: Cytoplasm (By similarity).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE=22829867; PubMed=12949112;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRAIN-W83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUCLEOTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=837;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Name=dnaE; OrderedLocusNames=PG0035; ORFNames=PG_0035;
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   UniProt
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   Consortium,
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Pred. No. 2e+03;
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
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   see http://www.uniprot.org/terms
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Best Local
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TIGR; PG0035; -
BioCyc; PG1N242619:PG0035-MONOMER; -
G0; G0:0005737; C:cytoplasm; IEA.
G0; G0:0008408; F:3'.5' exonuclease activ:
G0; G0:0003889; F:alpha DNA polymerase act
G0; G0:0003677; F:DNA binding; IEA.
G0; G0:0016740; F:transferase activity; II
G0; G0:0006260; P:DNA replication; IEA.
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Q8ILD1;
Q8ILD1;
01-MAR-2003,
01-MAR-2003,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-FEB-2006, e
Hypothetical p
ORFNames=PF14_
                                                                                                                                                                                                                                                                                   Gardner M.J., Hall N., Fung E., White O., Berriman M., Hyman R.W., Carlton J.M., Pain A., Nelson K.E., Bowman S., Paulsen I.T., James K., Eisen J.A., Rutherford K., Salzberg S.L., Craig A., Kyes S., Chan M.-S., Nene V., Shallom S.J., Suh B., Peterson J., Angiuoli S., Pertea M., Allen J., Selengut J., Haft D., Mather M.W., Vaidya A.B., Wartin D.M.A., Fairlamb A.H., Fraunholz M.J., Roos D.S., Ralph S.A., McFadden G.I., Cummings L.M., Subramanian G.M., Mungall C., Venter J.C., Carucci D.J., Hoffman S.L., Newbold C., Davis R.W., Fraser C.M., Barrell B.G.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      InterPro; IPR011708; UNA_pol3_alpha.
InterPro; IPR004365; OB tRNA_NA_bd.
InterPro; IPR003141; Pesterase_PHP_N.
InterPro; IPR004013; PHP_C.
InterPro; IPR004805; PolC_alpha.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TIGREAMs; TIGR00594; polc; 1.
Complete proteome; DNA replication; DNA-directed DNA polymerase;
Nucleotidyltransferase; Transferase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pfam; PF07733;
Pfam; PF02811;
                                     Hypothetical SEQUENCE 1:
                                                                                                    EMBL; AE014821; AAN36926.1; -;
InterPro; IPR002048; EF_hand_Ca
                                                                                                                                                               Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN=3D7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=36329;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Plasmodium falciparum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pfam; PF01336; tRNA_anti; 1.
SMART; SM00481; POLITIAC; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=22255705; PubMed=12368864; DOI=10.1038/nature01097;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUCLEOTIDE
                                                                        InterPro; IPR002048; EF_hand_Ca_bd.
PROSITE; PS00018; EF_HAND_1; UNKNOWN_1.
                                                                                                                                                                                                                               Nature 419:498-511(2002).
                                                                                                                                                                                                                                                     "Genome sequence falciparum.";
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PLAF7
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                                       al protein.
1267 AA;
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                                                                                                                                                                                                                                                                        of the
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85.7%;
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                                                                                                                                                                                                                                                                      human malaria parasite Plasmodium
                                     MW;
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Pred. No.
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exa; Haemosporida;
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a_bd.
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                                         239F91EBBD942C80 CRC64;
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2;
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Query Match

Similarity

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RESULT 53
PDS5_YEAST
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Matches
                                                                                                          MEDLINE=22923965; PubMed=14562106; DOI=10.1038/nature02046; Ghaemmaghami S., Huh W.-K., Bower K., Howson R.W., Belle A., Dephoure N., O'Shea E.K., Weissman J.S.; "Global analysis of protein expression in yeast."; Mature 425:737-741(2003).

-I-FUNCTION: Essential for the establishment and maintenance of sister chromatid cohesion at centromere proximal and distal regions during S phase. Also required for chromosomal
                                                                                                                                                                                                                                                                         involved
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FUNCTION, AND SUBCELLULAR LOCATION.

MEDLINE=20517447; PubMed=11062262; DOI=10.1083/jcb.151.3.613;

Hartman T., Stead K., Koshland D., Guacci V.,

"Pds5p is an essential chromosomal protein required for both sister chromatid cohesion and condensation in Saccharomyces cerevisiae.";

J. Cell Biol. 151:613-626(2000).
   Copyrighted
Distributed
                                             +
                                                                                                                                                                                                                                                                                                                PubMed=11864574; DOI=10.1016/S0960-9822(02)00681-4; Ivanov D., Schleiffer A., Eisenhaber F., Mechtler K
                                                                                                                                                                                                                                                                                                                                                                                          cohesion
                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=21065936; PubMed=11137006; DOI=10.1016/S0960-9822(00)00854-X; Panizza S., Tanaka T., Hochwagen A., Eisenhaber F., Nasmyth K.; Pds5 cooperates with cohesin in maintaining sister chromatid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "The nucleotide XIII.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bowman S., Churcher C.M., Badcock K., Brown D., Chillingworth T., Connor R., Dedman K., Devlin K., Gentles S., Hamlin N., Hunt S., Jagels K., Lye G., Moule S., Odell C., Pearson D., Rajandream M.A., Rice P., Skelton J., Walsh S.V., Whitehead S., Barrell B.G.; "The nucleotide sequence of Saccharomyces cerevisiae chromosome
            condensation.
I- INTERACTION:
Self; NbExp=1; IntAct=EBI-13077, E
FMM: Acetylated by ECO1.
-!- MISCELLANEOUS: Present with 7720 r
                                                                                                                                                                                                                                  LEVEL OF
                                                                                                                                                                                                                                                                                                       Nasmyth
                                                                                                                                                                                                                                                                                                                                                                              Curr. Biol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAIN=S288c / AB9
MEDLINE=97313268;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes; Saccharomycetales; Saccharomycetaceae; Saccharomyces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sisters protein 5).
Name=PDS5; OrderedLocusNames=YMR076C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            004264; Q04780;
01-NOV-1997, integrated into U
01-NOV-1997, sequence version
07-MAR-2006, entry version 35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Q04264;
                                                                                                                                                                                                                                                                                                                                                 ACETYLATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nature 387:90-93(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUCLEOTIDE SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=4932;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Saccharomyces cerevisiae (Baker's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sister chromatid
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Churcher C.M., Badcock
   by the UniProt Consortium, under the Creative Commons
                                                                                                                                                                                                                                                            12:323-328 (2002).
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            http://www.uniprot.org/terms
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A Lafontaine I., de Montigny J., Marck C., Neuveglise C., Talla E.,
A Lafontaine I., de Montigny J., Marck C., Neuveglise C., Talla E.,
A Goffard N., Frangeul L., Aigle M., Anthouard V., Babour A., Barbe V
A Barnay S., Blanchin S., Beckerich J.-M., Beyne E., Bleykasten C.,
A Boiszame A., Boyer J., Cattolico L., Confanioleri F., de Daruvar A.
Despons L., Fabre E., Fairhead C., Ferry-Dumazet H., Groppi A.,
A Hantraye F., Hennequin C., Jauniaux N., Joyet P., Kachouri R.,
A Hantraye F., Hennequin C., Jauniaux N., Joyet P., Kachouri R.,
A Nicaud J.-M., Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,
A Nicaud J.-M., Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,
A Pellenz S., Potter S., Richard G.-F., Straub M.-L., Suleau A.,
A Swennen D., Tekala F., Wesolowski-Louvel M., Westhof E., Witth B.,
A Swennen D., Tekala F., Wesolowski-Louvel M., Westhof E., Witth B.,
A Swennen D., Tekala F., Wesolowski-Louvel M., Westhof E., Witth B.,
A Swennen D., Tekala F., Wesolowski-Louvel M., Westhof E., Witth B.,
A Swennen D., Tekala F., Wesolowski-Louvel M., Westhof E., Witth B.,
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A Swennen D., Tekala F., Wesolowski-Louvel M., Westhof E., Witth B.,
A Swennen D., Tekala F., Wesolowski-Louvel M., Westhof E., Witth B.,
A Wicker P., Souciet J.-L.,
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19-JUL-2004,
21-FEB-2006,
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STRAIN=ATCC 2001 / C
PubMed=15229592; DOI
                             Copyrighted
Distributed
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GO; GO:0048002; F:protein self binding; IPI.
GO; GO:0007076; P:mitotic chromosome condensation; IMP.
GO; GO:000764; P:mitotic sister chromatid cohesion; IM
GO; GO:0006473; P:protein amino acid acetylation; IDA.
InterPro; IPR011989; ARM-like.
InterPro; IPR010357; HBAT.
Pfam; PF02985; HBAT; 1.
                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=5478;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Fungi; P
Saccharomycetales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Candida
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein
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Q6FNZ8;
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EMBL; Z48952; CAA88801.1;
PIR; S54451; S54451.
                                                                                Nature
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430:35-44(2004).
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CAG60997.1;
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DOI=10.1038/nature02579;
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                           Consortium, see http://www.uniprot.org/terms:ive Commons Attribution-NoDerivs License
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WA Hall N., Pain A., Berriman M., Churcher C.M., Harris B., Harris D.,

WA Mungall K.L., Bowman S., Atkin R., Baker S., Barron A., Brooks K.,

WA Buckee C.O., Burrows C., Cherevach I., Chillingworth C.,

Chillingworth T., Christodoulou Z., Clark L., Clark R., Corton C.,

Chillingworth T., Christodoulou Z., Clark L., Clark R., A Coronin A., Davies R.M., Davis P., Dear P., Dearden F., Doggett J.,

A Cronin A., Davies R.M., Davis P., Dear P., Dearden F., Doggett J.,

A Feltwell T., Goble A., Goodhead I., Gwilliam R., Hamlin N., Hance Z.,

A Harper D., Hauser H., Hornsby T., Holroyd S., Horrocks P.,

A Humphray S., Jagels K., James K.D., Johnson D., Kerhornou A.,

A Humphray S., Jagels K., James K.D., Johnson D., Lemnard N.,

A Knights A., Konfortov B., Kyes S., Larke N., Lawson D., Lemnard N.,

A Knights A., Konfortov B., Kyes S., Larke N., Lawson D., Lemnard N.,

A Knights A., Maddison M., Mclean J., Mooney P., Moule S., Murphy L.,

A Cliver K., Ormond D., Price C., Quail M.A., Rabbinowitsch E.,

A Cliver K., Sharp S., Smith R., Squares R., Squares S., Stevens K.,

A Seeger K., Sharp S., Smith R., Squares R., Squares S., Stevens K.,

A Sulston J.E., Craig A., Newbold C., Barrell B.G.;

"Sequence of Plasmodium falciparum chromosomes 1, 3-9 and 13.";

"Sequence of Plasmodium falciparum chromosomes 1, 3-9 and 13.";
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GO; GO:0016020; C:membrane; IEA.
GO; GO:0030870; C:Mre1l complex; IEA.
GO; GO:0005524; F:ATP binding; IEA.
GO; GO:0005524; F:ATP binding; IEA.
GO; GO:000587; F:ATPase activity; IEA.
GO; GO:0008270; F:zinc ion binding; IEA.
GO; GO:0008270; F:zinc ion binding; IEA.
GO; GO:0051276; P:chromosome organizati
GO; GO:00051276; P:chromosome organizati
GO; GO:0006810; P:transport; IEA.
InterPro; IPR003439; ABC_transp_like.
InterPro; IPR003439; ABC_transp_like.
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01-MAR-2003,
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Eukaryota; Alveolata;
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Hypothetical protein PFI0185w.
                                                                                       InterPro; IPR013320; ConA_like_subgrp.
InterPro; IPR004043; LCCL.
InterPro; IPR000772; Ricin_B_lectin.
                                                                                                                                                                                    EMBL; AL929355; CAD51723.1;
                                                                                                                                                                                                                                               Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                              Copyrighted
                                                                                                                                                                                                                                                                                                                                         Nature 419:527-531(2002).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=36329;
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m; PF04423; Rad50 zn_hook; 1.
n; PF02463; Rad50 zn_hook; 1.
n; PF02463; SMC_N; 1.
PAMS; TIGROUGG
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                                                                 PF03815; LCCL;
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PS50231; RICIN_B_LECTIN;
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                                                                                                                                                                                                                                                                              by the UniProt Consortium,
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F:zinc ion binding; IEA.
P:chromosome organization and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C:membrane; IEA.
C:Mrell complex; IEA.
F:ATP binding; IEA.
F:ATPase activity; IEA
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Pred. No. 2.2e+03;
Pred. No. 2.1e+03;
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RESULT
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Matches 6
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Submitted (SEP-2007)
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QBIBA8;
01-MAR-2003,
                                                                                                                                     05-JUL-2005, integrated into UniProtKB/TrEMBL.
05-JUL-2005, sequence version 1.
07-FBB-2006, entry version 4.
Hypothetical protein (Fragment).
ORFNames=PC00259,03.0;
                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL; AL844507; CAD51099.1; -; Genom GO; GG:0003723; F:RNA binding; IEA. InterPro; IPR000804; Clat adaptor s. InterPro; IPR012340; OB NĀ bd sub. InterPro; IPR013029; SI_RNĀ_bd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hypothetical protein; Repeat. seQUENCE 1523 AA; 178335 MW;
NUCLEOTIDE SEQUENCE:
PubMed=1563727; DOI=10.1126/science.1103717;
PubMed=1563727; DOI=10.1126/science.1103717;
Hall N., Karras M., Raine J.D., Carlton J.M., Kooij T.W.A.,
Berriman M., Florens L., Janssen C.S., Pain A., Christophides
Berriman M., Florens L., Janssen C.S., Harris D., Churcher C.M.,
James K., Rutherford K., Harris B., Harris D., Churcher C.M.,
Quail M.A., Ormond D., Doggett J., Trueman H.E., Mendoza J.,
Bidwell S.L., Rajandream M.A., Carucci D.J., Yates J.R. III,
                                                                                                                                                                                                   Q4XUN1 PLACH
Q4XUN1;
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Eukaryota; Alveolata; Apicomplexa;
                                                                                                                                                                                                                                                                                                                                                                                Hypothetical SEQUENCE 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
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                                                                                                              Plasmodium chabaudi.
Eukaryota; Alveolata;
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PROSTTE; PS00989; CLAT_ADAPTOR_S; UNKNOWN_1.
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                                                                                                 NCBI_TaxID=5825;
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Pred. No. 2.8e+0
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                                     Kooij T.W.A.,
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Hypothetical
SEQUENCE 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-JUL-2005,
05-JUL-2005,
07-FEB-2006,
Hypothetical
                                                                                                                                                                                                                                                                                                                                                                   STRAIN-Ankara isolate clone C9;
PubMed=15994997; DOI=10.1258/jrsm.98.7.320;
Pain A., Renauld H., Berriman M., Murphy L., Yeats C.A., Weir W. Kerhornou A., Aslett M., Bishop R., Bouchier C., Cochet M., Coulson R.M.R., Cronin A., de Villiers E.P., Fraser A., Fosker N. Gardner M., Goble A., Griffiths-Jones S., Harris D.E., Katzer F. Gardner M., Goble A., Maser P., McKellar S., Mooney P., Morton F., Larke N., Lord A., Maser P., McKellar S., Mooney P., Morton F., Nene V., O'Neil S., Price C., Quail M.A., Rabbinowitsch E., Rawlings N.D., Rutter S., Saunders D., Seeger K., Shah T., Squar Squares S., Tivey A., Walker A.R., Woodward J., Dobbelaere D.A.E. Langsley G., Rajandream M-A., McKeever D., Shiels B., Tait A., Barrell B., Hall N.,
                                                                                                                                                                                                                                                Copyrighted Distributed
                                                                                                                                                                                                                                                                                                             Science
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUCLEOTIDE SEQUENCE
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Q4UBC4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       transcriptomic, and proteomic analyses.",
Science 307:82-86(2005)
-i- CAUTION: The sequence shown here is derived
EMBL/GenBank/DDBJ whole genome shotgun (WGS)
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155
                                                                                                                                                                                                     CR940352; CAI75877.1;
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                                                                              Similarity 6; Conserv
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                                                                                                 .3e+03;
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A Batzoglou S., Lee S.-I., Bastuerkmen M., Spevak C.C., Clutterbuck J.,

A Kapitonov V., Jurka J., Scazzocchio C., Farman M., Butler J.,

A Purcell S., Harris S., Braus G.H., Draht O., Busch S., D'Enfert C.,

A Bouchier C., Goldman G.H., Bell-Pedersen D., Griffiths-Jones S.,

A Doonan J.H., Yu J., Vienken K., Pain A., Freitag M., Selker E.U.,

A Doonan J.H., Penalva M.A., Oakley B.R., Momany M., Tanaka T.,

A Archer D.B., Penalva M.A., Oakley B.R., Momany M., Tanaka T.,

A Kumagai T., Asai K., Machida M., Nierman W.C., Denning D.W.,

Caddick M., Hynes M., Paoletti M., Fischer R., Miller B.L., Dyer P.S.,

Sachs M.S., Osmani S.A., Birren B.W.;

"Sequencing of Aspergillus nidulans and comparative analysis with A.

If fumigatus and A. oryzae.";

L. Nature 438:1105-1115(2005).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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Best Local (
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GO; GO:0005488; F:binding; IEA.
GO; GO:0008233; F:peptidase activity; IE
GO; GO:0051301; P:cell division; IEA.
GO; GO:0051301; P:cell division; IEA.
InterPro; IPR0105314; Peptidase C50.
InterPro; IPR010599; TPR-like helical.
PANTHER; PTHR12792; Peptidase C50; 1.
Pfan; PF03568; Peptidase C50; 1.
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P33144;
01-OCT-1993,
01-OCT-1993,
07-FEB-2006,
Separin (EC 3
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                 Emericella nidulans (Asperg
Eukaryota; Fungi; Ascomycot
Eurotiales; Trichocomaceae;
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26-APR-2005,
07-MAR-2006,
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Cell division;
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Eurotiales; Trichocomaceae; En
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                 nidulans (Aspergillus nidulans).
Fungi; Ascomycota; Pezizomycotina;
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Emericella.
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MEDLINE-99376085; PubMed=10448855; DOI=10.1038/22964;

MEDLINE-99376085; PubMed=10448855; DOI=10.1038/22964;

Bowman S., Lawson D., Basham D., Brown D., Chillingworth T.,

Churcher C.M., Craig A., Davies R.M., Devlin K., Feltwell T.,

Gentles S., Gwilliam R., Hamlin N., Harris D., Holroyd S., Hornsby

Horrocks P., Jagels K., Jassal B., Kyes S., McLean J., Moule S.,

Mungall K., Murphy L., Oliver K., Quail M.A., Rajandream M.-A.,

Rutter S., Skelton J., Squares R., Squares S., Sulston J.E.,

Whitehead S., Woodward J.R., Newbold C., Barrell B.G.;

"The complete nucleotide sequence of chromosome 3 of Plasmodium
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MEROP8; C50.001; -...
InterPro; IPR005314; Peptidase C50.
InterPro; IPR011990; TPR-like helical.
PANTHER; PTHR12792; Peptidase C50; 1.
PANTHER; PTHR12792; Peptidase C50; 1.
Cell cycle; Cell division; Chromosome partition;
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MEDLINE=92348436; PubMed=1639810;
May G.S., McGoldrick C.A., Holt C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-MAR-2003, sequence version 07-FEB-2006, entry version 21 Hypothetical protein MAL3P2.2
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[2]
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               MEDLINE=22255708; Pul
Hall N., Pain A., Ber
Mungall K.L., Bowman
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NCBI_TaxID=36329;
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SIMILARITY: Belongs to the peptidase C50 family.
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FUNCTION: Required for nuclear division. Could function in the mitotic spindle.

mitotic spindle.

CATALYTIC ACTIVITY: All bonds known to be hydrolyzed by this endopeptidase have arginine in P1 and an acidic residue in P4. P6 is often occupied by an acidic residue or by an hydroxy-amino-acid residue, the phosphorylation of which enhances cleavage.
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1 2067 Separin.
/FTId=PRO_0000205902
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DYKDDDD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1964
2067
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synonyms=PFC0165w;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1964
IA; 227926
                   PubMed=12368867; DOI=10.1038/nature01095;
Berriman M., Churcher C.M., Harris B., Ha
man S., Atkin R., Baker S., Barron A., Bro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            [GENOMIC DNA]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              80.9%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (isolate 3D7)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Apicomplexa; Haemosporida; Plasmodium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chromosome partition; Hydrolase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    By similarity. MW; E0655D939EC148DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 38;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                UniProtKB/TrEMBL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ω.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2226
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 2067;
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s B., s.c.
A., Brooks
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                                                              Harris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mitosis
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Cronin A., Davies R.M., Davis P., Dear P., Dearden F., Doggett J.,
A. Feltwell T., Goble A., Goodhead I., Gwilliam R., Hamlin N., Hance Z.,
A. Harper D., Hauser H., Hornsby T., Holroyd S., Horrocks P.,
A. Humphray S., Jagels K., James K.D., Johnson D., Kerhornou A.,
A. Knights A., Konfortov B., Kyes S., Larke N., Lawson D., Lennard N.,
A. Line A., Maddison M., Mclean J., Mooney P., Moule S., Murphy L.,
A. Oliver K., Ormond D., Price C., Quall M.A., Rabbinowitsch E.,
A. Oliver K., Ormond D., Price C., Quall M.A., Rabbinowitsch E.,
A. A. Rajandream M.A., Rutter S., Rutherford K.M., Sanders M., Simmonds
A. Seeger K., Sharp S., Smith R., Squares R., Squares S., Stevens K.,
A. Seeger K., Tivey A., Unwin L., Whitehead S., Woodward J.,
A. Taylor K., Tivey A., Newbold C., Barrell B.G.;

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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                          PubMed=1553466; DOI=10.1126/science.1103773;
Loftus B.J., Fung E., Roncaglia P., Rowley D., Amedeo P., Bruno D.,
Vamathevan J., Miranda M., Anderson I.J., Fraser J.A., Allen J.E.,
Bosdet I.E., Brent M.R., Chiu R., Doering T.L., Donlin M.J.,
Bosdet I.E., Brent M.R., Chiu R., Doering T.L., Donlin M.J.,
Bosdet I.E., Brent M.R., Chiu R., Doering T.L., Donlin M.J.,
Huang J.C., Janbon G., Jones S.J.M., Koo H.L., Krzywinski M.I.,
Kwon-Chung K.J., Lengeler K.B., Maiti R., Marra M.A., Marra R.E.,
Mathewson C.A., Mitchell T.G., Pertea M., Riggs F.R., Salzberg S.L.,
Schein J.E., Shvartsbeyn A., Shin H., Shumway M., Specht C.A.,
Suh B.B., Tenney A., Utterback T.R., Wickes B.L., Wortman J.R.,
Wye N.H., Kronstad J.W., Lodge J.K., Heitman J., Davis R.W.,
Fraser C.M., Hyman R.W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 D7_CRYNE
Q5KID7_C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Fungi; Basidiomycota; Hymenomycetes;
Tremellomycetidae; Tremellales; Tremellaceae; Fi
NCBI_TaxID=5207;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-FEB-2006, entry version 9.
1-phosphatidylinositol-3-phosphate
OrderedLocusNames=CND03330;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; AL034558; CAB38989.3; -; Genomic_DNA. GO; GO:0005815; C:microtubule organizing ce GO; GO:000922; C:spindle pole; IEA. GO; GO:0000226; P:microtubule cytoskeleton
                                                                                        Fraser C.M., Hyman R.W.;
"The genome of the basidionycetous
Cryptococcus neoformans.";
Science 307:1321-1324(2005).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Q5KID7;
15-FEB-2005,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pfam; PF04130; Spc97
Hypothetical protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    InterPrc; IPR007259; Spc97_Spc98.
PANTHER; PTHR19302; Spc97_Spc98; 1.
Pfam; PF04130; Spc97_Spc98; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Copyrighted
Distributed
                      Copyrighted Distributed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cryptococcus neoformans (Filobasidiella neoformans)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-FEB-2005, sequence version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    218
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6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
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                      by the under t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    integrated into UniProtKB/TrEMBI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AA;
                        the
                                              UniProt Consortium,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                [LARGE SCALE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          80.9%;
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                           Creative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cherevach I., Chillingworth odoulou Z., Clark L., Clark I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 38; DB Pred. No. 3.8e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
                        sortium, see http://www.uniprot.org/terms
Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENOMIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8690501ED4994768 CRC64;
                                                                                                                                            yeast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5-kinase,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2384
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                                                                                                                                              and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA]
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                                                                                                                                            human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Heterobasidiomycetes;
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( GO; GO:0016308; F:1-phosphatidylinositol-4-phosphate 5-kinase...

( GO; GO:0005524; F:ATP binding; IEA.

( GO; GO:0005521; F:protein binding; IEA.

( R GO; GO:0005215; F:protein binding; IEA.

( R GO; GO:0008270; F:zinc ion binding; IEA.

( R GO; GO:004267; P:cellular protein metabolism; IEA.

( R GO; GO:0044267; P:cellular protein metabolism; IEA.

( R InterPro; IPR002423; Cpn60/TCP-1.

( R InterPro; IPR002498; PIPSK.

( R InterPro; IPR00306; Znf FYVE.

( Pfam; PF01363; FYVE; I.

( DR SMART; SM00064; FYVE; I.

DR PROSITE; PS50178; ZF FYVE; I.

DR PROSITE; PS50178; ZF FYVE; I.
    Query Match
Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 6
                                                                          InterPro; IPR002498; PIPSK.
InterPro; IPR000306; Znf FYVE.
Pfam; PF00118; Cpn60 TCPI; 1.
Pfam; PF001363; FYVE; 1.
Pfam; PF01504; PIPSK; 1.
SMART; SM00064; FYVE; 1.
PROSITE; PS50178; ZF FYVE; 1.
                                                                                                                                                                  GG; GG:0016308; F:1-phosphatidylinositol-4-phosphate GO; GG:00015524; F:ATP binding; IEA.
GG; GG:00015515; F:protein binding; IEA.
GG; GG:0008270; F:zinc ion binding; IEA.
GG; GG:0044267; F:cellular protein metabolism; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Q55U02
Q55U02
                                                                                                                                                                                                                                                                                                               "Cryptococcus neoformans serotype D sequencing."
Submitted (JUL-2004) to the EMBL/GenBank/DDBJ dat
-I- CAUTION: The sequence shown here is derived:
EMBL/GenBank/DDBJ whole genome shotgun (WGS)
                                                                                                                                                                                                                                                                                                                                                                                                                                              Cryptococcus neoformans var. neoformans B-3501A.

Eukaryota; Fungi; Basidiomycota; Hymenomycetes; Heterobasidiomycetes;

Tremellomycetidae; Tremellales; Tremellaceae; Filobasidiella.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hypothetical protein. ORFNames=CNBD3020;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24-MAY-2005,
07-FEB-2006,
                                                      Hypothetical SEQUENCE 2
                                                                                                                                                                                                                                      EMBL; AAEY01000020; EAL21247.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                              Distributed under the Creative Commons
                                                                                                                                                                                                                                                                            Copyrighted by the UniProt Consortium,
                                                                                                                                                                                                                                                                                                                                                                    Fung E., Hyman R.W., Rowle Wickes B.L., Fu J., Davis
                                                                                                                                                                                                                                                                                                                                                                                             STRAIN=B-3501A;
                                                                                                                                                                                                                                                                                                                                                                                                          NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=283643;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-MAY-2005,
                                                                                                                                                                                                                                                                                                   preliminary data.
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    Similarity 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DYKDDDD
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                                                      il protein.
2432 AA;
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  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence version entry version 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          integrated into
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRELIMINARY;
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             80.9%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                    Rowley D.,
Davis R.W.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 80.9%;
85.7%;
                                                      268925
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        263686 MW;
 Score 38; DB
Pred. No. 4.2e
1; Mismatches
                                                      MW.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 38;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          UniProtKB/TrEMBL
                                                                                                                                                                                                                                                                                                                                                                                Bruno D.,
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                                                      A5A9E486EDC898EE CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AD29D2B0BF62F793 CRC64;
                                                                                                                                                                                                                                                                                                                                                      sequencing.";
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01-JUN-2002,
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                    Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                           NUCLEOTIDE SEQUENCE.
Fowler E., Peters J., Gatton M., Chen N., Cheng Q., Submitted (AUG-2001) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                  NUCLEOTIDE SEQUENCE.
MEDLINE=21839615; PubMed=11849711;
Fowler E.V., Peters J.M., Gatton M.
"Genetic diversity of the DBLalpha
                                                                                                                                                                                                                                                                                                                         Plasmodium falciparum
Eukaryota; Alveolata;
NCBI_TaxID=5833;
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Seeger K., Murphy L., Harris D., Berriman M., Pain A., H
Quail M., Barrell B.;
Submitted (SEP-2002) to the EMBL/GenBank/DDBJ databases.
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Eukaryota; Alveolata;
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GO:0004222; F:metalloendopeptidase
GO:0009405; P:pathogenesis; IEA.
GO:0006508; P:proteolysis; IEA.
GO:011591; Botulinum.
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Biochem. Parasitol. 120:117-126(2002)
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.L., Chen N., Cheng Q.;
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n Plasmodium
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MEDLINE=22255706; PubMed=12368865; DOI=10.1038/nature01099; Carlton J.M., Angiuoli S.V., Suh B.B., Kooij T.W., Pertea M., Silva J.C., Ermolaeva M.D., Allen J.E., Selengut J.D., Koo H.L. Peterson J.D., Pop M., Kosack D.S., Shumway M.F., Bidwell S.L., Shallom S.J., van Aken S.E., Riedmuller S.B., Feldblyum T.V.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL;
                                                                                                                                                                       ORFNames=PY06254;
Plasmodium yoelii yoelii.
Bukaryota; Alveolata; Apicomplexa; Haemosporida;
                                                                                                                                                                                                                                     15-DEC-2003, integrated into UniProtKB/TrEMBL
15-DEC-2003, sequence version 1.
07-FEB-2006, entry version 11.
Hypothetical protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PubMed=1512253; DOI=10.1086/383250;
Kaestli M., Cortes A., Lagog M., Ott M., Beck H.-P
"Longitudinal Assessment of Plasmodium falciparum
Transcription in Naturally Infected Asymptomatic C
                                                                                                                                                                                                                                                                                                               Q7RB94_PLAYO
Q7RB94;
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                                                                                                                      NUCLEOTIDE
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                                                                                                                                                                                                                                                              PubMed=15637271; DOI=10.1126/science.1103717;
Hall N., Karras M., Raine J.D., Carlton J.M., Kooij T.W.A.,
Hall N., Karras M., Raine J.D., Carlton J.M., Kooij T.W.A.,
Harris D., Christophides G.K.,
James K., Rutherford K., Harris B., Harris D., Churcher C.M.,
James K., Rutherford K., Harris B., Harris D., Churcher C.M.,
Quail M.A., Ormond D., Doggett J., Trueman H.E., Mendoza J.,
Bidwell S.L., Rajandream M.A., Carucci D.J., Yates J.R. III,
Kafatos F.C., Janse C.J., Barrell B.G., Turner C.M.R., Waters A.P.,
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Nature 419:512-519 (2002).
                                                                                                                                                                                                Plasmodium chabaudi.
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                                                                                             Hypothetical SEQUENCE 1
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                                                                                                                                                                                                                                                       Sinden R.S.;
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n L.W., Vaidya A.B.,
, Smith H.O., White O.R.,
M., Hoffman S.L., Gardner M.J.,
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       NUCLEOTIDE SEQUENCE.
PubMed=15637271; DOI=
Hall N., Karras M., F
Berriman M., Florens
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Chow T.-Y., Hsing Y.-I.C., Chen C.-S., Chen H.-H., Liu S.-M., Choo Y.-T., Lee P.-F., Chang S.-J., Chen H.-C., Chen S.-K., Chen T.-R., Chen Y.-L., Cheng C.-H., Chung C.-I., Han S.-Y., Chen T.-R., Chen Y.-L., Cheng C.-H., Chung C.-I., Han S.-Y., Chen T.-R., Hsiung J.-N., Hsu C.-H., Hsu C.-T., Kau P.-I., Lee Lee M.-C., Leu H.-L., Li Y.-F., Lin S.-J., Lin Y.-C., Lu P.-C., Wu S.-W., Yang K.-C., Yu C.-Y., Yu S.-W., Wu H.-P., Shaw J.-F., Wu S.-W., Yang K.-C., Yu C.-Y., Yu S.-W., "Oryza sativa BAC OSUNBOO86G17 genomic sequence.";
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Chow Y.-Y., Hsing Y.-I.C., Chen C.-S., Chen H.-H., Liu S.-M., Chon Y.-T., Chang S.-J., Chen H.-C., Chen S.-K., Chen T.-R., Chen Y.-L., Chang C.-H., Chung C.-I., Han S.-Y., Hsiao S.-H., Hsiung J.-N., Hsu C.-H., Huang J.-J., Kau P.-I., Lee M.-C., L Hsiung J.-N., Lin S.-J., Lin Y.-C., Wu H.-P., Shaw J.-F.;

"Oryza sativa BAC OSJNBA0030114 genomic sequence.";
                                                                                                                                      Plasmodium chabaudi.
Eukaryota; Alveolata;
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NCBI TaxID=39947;
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AAV59389.1;
DOI=10.1126/science.1103717;
M., Raine J.D., Carlton J.M., K
rens L., Janssen C.S., Pain A.,
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QZZA91;
Q2ZA91;
20-DEC-2005, integrated into
20-DEC-2005, sequence version
07-FEB-2006, entry version 3.
Hypothetical protein.
ORFNames=ShewmrdDRAFT_1640;
                                                                                                                                                                                                                                                                                     Larimer F., Land M.;

"Annotation of the draft genome of Shewanella sp. MR-4.";

"Bubmitted (NOV-2005) to the EMBL/GenBank/DDBJ databases.
-!- CAUTION: The sequence shown here is derived from an
-!- EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which
                                                                                                            Hypothetical SEQUENCE 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRAIN=MR-4;
US DOE Joint Genome Institute (JGI-PGF);
Copeland A., Lucas S., Lapidus A., Barry K., I
Copeland A., Incas S., Pitluck S., Richardson E
Hammon N., Israni S., Pitluck S., Richardson E
"Sequencing of the draft genome and assembly c
Submitted (OCT-2005) to the EMBL/GenBank/DDBJ
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Quail M.A., Ormond D., Doggett J., Truema
Bidwell S.L., Rajandream M.A., Carucci D.,
Kafatos F.C., Janse C.J., Barrell B.G., T
Sinden R.S.,
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Shewanellaceae; Shewanella.
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SEQUENCE 168 AA; 2
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EMBL/GenBank/DDBJ whole
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J., Trueman H.E., Mendoza J.,
Carucci D.J., Yates J.R. III,
ell B.G., Turner C.M.R., Water
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Attribution-NoDerivs License
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Q35Y65, 9GAMM
PRELIMINARY; Pl
Q35Y65, 1000
06-DEC-2005, integrated into Un
06-DEC-2005, sequence version 1
07-FEB-2006, entry version 3.
Hypothetical protein.
ORFNames=Shewmr7DRAFT_1872;
                                                                                                                                                                                                         Q366F4;
06-DE~
                                                                                                                                                 06-DEC-2005, integrated into U
06-DEC-2005, sequence version
07-PEB-2006, entry version 3.
Hypothetical protein.
ORFNames=Shewana3DRAFT_0257;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Larimer F., Land M.;

"Annotation of the draft genome of Shewanella sp. MR-7.";

"Annotation of the EMBL/GenBank/DDBJ databases.

Submitted (NOV-2005) to the EMBL/GenBank/DDBJ databases.

-I- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US DOE Joint Genome Institute (JGI-PGF);
Copeland A., Lucas S., Lapidus A., Barry K., Detter J.C., G
Hammon N., Israni S., Pitluck S., Richardson P.;
"Sequencing of the draft genome and assembly of Shewanella
Submitted (OCT-2005) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                     Hypothetical SEQUENCE 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Copyrighted by the UniProt Consortium, Distributed under the Creative Commons
            US DOE Joint Genome Institute (JGI-PGF);
COpeland A., Lucas S., Lapidus A., Barry K., Detter J.C., Glavina Hammon N., Israni S., Pitluck S., Richardson P.;
"Sequencing of the draft genome and assembly of Shewanella sp. ANJ
                                                                                                                Bacteria; Proteobacteria; Gammaproteobacteria; Shewanellaceae; Shewanella.
                                                                                                                                           Shewanella sp. ANA-3.
                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL;
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  "Sequencing 3.";
                                                                                                    NCBI_TaxID=94122;
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85.7%;
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Pred. No. 3.8e+02;
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Q8EFW6;
01-MAR-2003,
01-MAR-2003,
                                                                                                                                                                                                                    MEDINE=22297686; PubMed=12368813; DOI=10.1038/nbt749; MEDINE=22297686; Paulsen I.T., Nelson K.E., Gaidos E.J., Nelson W.C. Heidelberg J.F., Paulsen I.T., Nelson K.E., Gaidos E.J., Nelson W.C. Read T.D., Eisen J.A., Seshadri R., Ward N.L., Methe B.A., Clayton R.A., Meyer T., Tsapin A., Scott J., Beanan M.J., Durkin A.S., Brinkac L.M., Daugherty S.C., DeBoy R.T., Dodson R.J., Durkin A.S., Haft D.H., Kolonay J.F., Madupu R., Peterson J.D., Umayam L.A., White O., Wolf A.M., Vamathevan J.J., Weidman J.F., Impraim M., White O., Wolf A.M., Vamathevan J.J., Weidman J.F., Impraim M., Lee K., Berry K.J., Lee C., Mueller J., Khouri H.M., Gill J., Utterback T.R., McDonald L.A., Feldblyum T.V., Smith H.O., Venter J.C., Nealson K.H., Fraser C.M.; "Genome sequence of the dissimilatory metal ion-reducing bacterium "Genome sequence of the dissimilatory metal ion-reducing bacterium"
                                                                                      BioCyc; S
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SEQUENCE
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Hypothetical protein.
ORFNames=SO_1849;
Shewanella oneidensis.
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US DOE Joint Genome Institute (JGI-ORNL);
Larimer F., Land M.;
"Annotation of the draft genome of Shewanella sp. ANA-3."
Submitted (NOV-2005) to the EMBL/GenBank/DDBJ databases.
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                                                                                                                                                        Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                              Shewanella oneidensis.";
Nat. Biotechnol. 20:1118-1123(2002).
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Commons Attribution-NoDerivs License
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Hypothetical protein (Frc
ORFNames=PB000975.00.0;
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                                                                                                                                                Copyrighted Distributed
                                                                                                                                                                                                                                                                     transcriptomic, and proteomic Science 307:82-86(2005).
                                                                Hypothetical NON_TER
                                                                                                                                                                                                                                                                                                              "A comprehensive survey of the Plasmodium
                                                                                                                                                                                                                                                                                                                                      Sinden R.S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUCLEOTIDE SEQUENCE
PubMed=15637271; DOI
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Eukaryota; Alveolata;
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                                                                                                                                                                                                                                                         05-JUL-2004,
05-JUL-2004,
07-MAR-2006,
Hypothetical
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Hypothetical pr
ORFNames=cgd5_4
                                                                                                                                                                                                                                                                                                                                          Q74HZ2_
Q74HZ2;
                                                                    NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNAJ
STRAIN-NCC 533;
PubMed=14983040; DOI=10.1073/pnas.0307327101;
Pridmore R.D., Berger B., Desiere F., Vilanova D., Barretto C., Pittet A.-C., Zwahlen M.-C., Rouvet M., Altermann E., Barrangou Mollet B., Mercenier A., Klaenhammer T., Arigoni F., Schell M.A. "The genome sequence of the probiotic intestinal bacterium
                                                                                                                                                                                                                                         OrderedLocusNames=LJ0673;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL; AAEE01000017; EAK87298.1; -;
Hypothetical protein.
SEQUENCE 259 AA; 29923 MW; E37E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PubMed=15044751; DOI=10.1126/science.1094786; Abrahante Abrahamsen M.S., Templeton T.J., Enomoto S., Abrahante Lancto C.A., Deng M., Liu C., Widmer G., Tzipori S., Bu Bankler A.T., Dear P.H., Konfortov B.A., Spriggs H.F., Anantharaman V., Aravind L., Kapur V.; "Complete genome sequence of the apicomplexan, Cryptosp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cryptosporidium parvum.
Eukaryota; Alveolata; Apicomplexa;
Cryptosporidiidae; Cryptosporidium.
                                                                                                                                                                                                  Bacteria; Firmicutes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Copyrighted by the UniProt Consortium, see http://www.uniprot.org/to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRAIN=Iowa type PubMed=15044751;
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Q5CPE3;
                                                                                                                                                          NCBI_TaxID=33959;
                                                                                                                                                                                   Lactobacillus
                                                                                                                                                                                                                      Lactobacillus johnsoni:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Science 304:441-445(2004).
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RA Eichinger L., Pachebat J.A., Gloeckner G., Rajandream M.A.,
RA Eichinger L., Pachebat J.A., Gloeckner G., Szafranski K., Xu Q.,
RA Sucgang R., Berriman M., Song J., Olsen R., Szafranski K., Xu Q.,
RA Sucgang R., Berriman M., Song J., Olsen R., Szafranski K., Xu Q.,
RA Sucgang R., Berriman M., Song J., Olsen R., Rivero F.,
RA Tunggal B., Kummerfeld S., Madera M., Konfortov B.A., Rivero F.,
RA Pilcher K., Chen G., Saunders D., Sodergren E.J., Davis P.,
RA Pilcher K., Chen G., Saunders D., Sodergren E.J., Davis P.,
RA Pilcher K., Nie X., Hall N., Anjard C., Hemphill L., Bason N.,
RA Kerhornou A., Nie X., Hall N., Anjard C., Hemphill L., Bason N.,
RA Farbrother P., Desany B., Just E., Morio T., Rost R., Churcher C.M.,
RA Cooper J., Haydock S., van Driessche M., Cronin A., Goodhead I.,
RA Cooper J., Haydock S., van Driessche M., Kaprer D., Lindsay R.,
RA Muzny D.M., Mourier T., Pain A., Lu M., Harper D., Lindsay R.,
RA Muzny D.M., Mourier T., Pain A., Lu M., Harper D., Lindsay R.,
RA Hauser H., James K.D., Quiles M., Madan Babu M., Saito T.,
RA Hauser H., James K.D., Quiles M., Madan Babu M., Saito T.,
RA Hauser H., James K.D., Quiles M., Mangavelu M., Johnson D.,
RA Knights A., Loulseged H., Mungall K.L., Oliver K., Price C.,
RA Misha A., Urushihara H., Hernandez J., Rabbinowitsch E., Steffen D.,
RA Guail M.A., Urushihara H., Hernandez J., Rabbinowitsch E., Steffen D.,
RA Tivey A., Sugano S., White B., Walker D., Woodward J.R., Winckler T.,
RA Tanaka Y., Shaulsky G., Schleicher M., Weinstock G.M., Rosenthal A.,
RA Kuena A., Williams J.G., Dear P.H., Noegel A.A., Barrell B.G.,
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24-MAY-2005, sequence version 1.
07-FBB-2006, entry version 3.
Hypothetical protein.
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GenomeReviews; AE017198 GR; I.00673.
BioCyc; L.004257314:LJ0673-MONOMER; -.
GO; GO:0003824; F:catalytic activity; IEA.
Hypothetical SEQUENCE 2
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ORFNames=DDB0204132;
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InterPro; IPR000379; Ser_estrs.
Pfam; PF07859; Abhydrolase_3; 1.
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                                                                      EMBL; AAFI01000072; EAL66940.1; -; Genomic_DNA
                                                                                                                                                Distributed
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   BEA0D00562DC563E CRC64;
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                                                                                                                                                                                                                                                                                                                                   RC STRAIN-927/4 GUTatiol.;

RX PubMed=16020726; DOI=10.1126/science.1112642;

RX Berriman M., Ghedin E., Hertz-Fowler C., Blandin G., Renauld H.,

RA Bartholomeu D.C., Lennard N.J., Caler E., Hamlin N.E., Haas B.,

RA Bohme U., Hannick L., Aslett M.A., Shallom J., Marcello L., Hou L.,

RA Bingaud F., Brooks K., Carrington M., Cherevach I.,

RA Wickstead B., Alsmark U.C.M., Arrowsmith C., Atkin R.J., Barron A.J.,

RA Bringaud F., Brooks K., Carrington M., Cherevach I.,

RA Bringaud F., Brooks K., Carrington M., Cherevach I.,

RA Hingworth T.-J., Churcher C., Clark L.N., Corton C.H., Cronin A.,

RA All Chillingworth T.-J., Churcher C., Clark L.N., Corton C.H., Cronin A.,

RA All Chillingworth T.-J., Churcher C., Clark L.N., Corton C.H., Cronin A.,

RA Chillingworth T.-J., Djikeng A., Feldbyum T., Field M.C.,

RA Chillingworth T.-J., Djikeng A., Feldbyum T., Field M.C.,

RA Hostetler J., Tvens A., Jagels K., Johnson D., Johnson J., Jones K.,

RA Hostetler J., Ivens A., Jagels K., Johnson D., Johnson J., Jones K.,

RA Kerhornou A.X., Koo H., Larke N., Landfear S., Larkin C., Leech V.,

RA Hostetler J., Mungall K., Norbertczak H., Ormond D., Pai G.,

RA Morgan G.W., Mungall K., Norbertczak H., Ormond D., Pai G.,

RA Rajandream M.-A., Reitter C., Salzberg S.L., Sanders M., Schobel S.,

RA Rajandream M.-A., Reitter C., Salzberg S.L., Sanders M., Schobel S.,

RA Rajandream M.-A., Namerell B.G., Donelson A.J., Tallon L., Turner C.M.R.,

RA Wilte B., White O., Whitehead S., Woodward J., Wortman J., Adams M.D.,

Rambley T.M., Gull K., Ullu E., Barry J.D., Fairlamb A.H.,

Opperdoes F., Barrell B.G., Donelson J.E., Hall N., Fraser C.M.,

"The genome of the African trypanosome Trypanosoma brucei.";

RI Science 300-416-427(2015)
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PubMed=16020724; DOI=10.1126/science.1112181;

El-Sayed N.M.A., Wyler P.J., Blandin G., Berriman M., Crabtree J. Aggarwal G., Caler E., Renauld H., Worthey E.A., Hertz-Powler C., Aggarwal G., Caler E., Bartholomeu D.C., Haas B.J., Tran A.-N., Mortman J.R., Alsmark U.C.M., Angiuoli S., Anupama A., Badger J., Bringaud F., Cadag E., Carlton J.M., Cerqueira G.C., Creasy T., Bringaud F., Cadag E., Carlton J.M., Hauser C., Ivens A.C., Kummerfeld S.K., Pereira-Leal J.B., Nilsson D., Peterson J., Salzberg S.L., Shallom J., Silva J.C., Sundaram J., Westenberger Mhite O., Melville S.E., Donelson J.E., Andersson B., Stuart K.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hypothetical protein. ORFNames=Tb09.211.2220;
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Distributed
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                                                                                                  38122
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Pred.
   Score 37; Pred. No.
                                                                                                  AB5C219A514BA4B0 CRC64;
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, Hertz-Fowler C.,
Tran A.-N.,
                                Length 335
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7.4e+02;

NUCLEOTIDE SEQUENCE [MRNA].

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01-JAN-1998, sequence version 1.

07-MAR-2006, entry version 30.

Bromelain precursor (EC 3.4.22.32) (Allergen Ana c 2).

Bromelain precursor (EC 3.4.22.32) (Allergen Ana c 2).

Ananas comosus (Pineapple).

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Eukaryota; Viridiplantae; Streptophyta; Doales; Bromeliaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dujon B., Sherman D., Fischer G., Durrens P., Casaregola S.,
Lafontaine I., de Montigny J., Marck C., Neuveglise C., Talla E.,
Goffard N., Frangeul L., Aigle M., Anthouard V., Babour A., Barbe V.,
Barnay S., Blanchin S., Beckerich J.-M., Beyne E., Bleykasten C.,
Bolsrame A., Boyer J., Cattolico L., Confanioleri F., de Daruvar A.,
Despons L., Fabre E., Fairhead C., Ferry-Dumazet H., Groppi A.,
Hantraye F., Hennequin C., Jauniaux N., Joyet P., Kachouri R.,
Kerrest A., Koszul R., Lemaire M., Lesur I., Maller H.,
Nicaud J.-M., Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,
Pellenz S., Potier S., Richard G.-F., Straub M.-L., Suleau A.,
Swennen D., Tekaia F., Wesolowski-Louvel M., Westhof E., Witth B.,
Zeniou Mever M., Tekaia F., Wesolowski-Louvel M., Westhof E., Witth B.,
Zeniou Mever M., Tivanovic V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-AUG-2004,
16-AUG-2004,
07-FEB-2006,
  NCBI_TaxID=4615;
                                                                                                                                                                                                                                                                                                      Complete proteome. SEQUENCE 341 AA;
                                                                                                                                                                                                                                                                                                                                                              Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC STRAIN-CLIB 122 / E 150; PubMed=15229592; DOI=10.1038/nature02579; Dujon B., Sherman D., Fischer G., Durrens
                                                                                                                                                                                                                                                                                                                               InterPro; IPR013256; Chromatin
                                                                                                                                                                                                                                                                                                                                                                                                          Nature 430:35-44(2004).
                                                                                                                                                                                                                                                                                                                                                                                                                                  Bouchier C., Caudron B., Scarpelli C., Wincker P., Souciet J.-L.;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Zeniou-Meyer M., Zivanovic Y., Bolotin-Fukuhara M., Thierry A., Bouchier C., Caudron B., Scarpelli C., Gaillardin C., Weissenba
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            lipolytica.
OrderedLocusNames=YALIOD08734g;
OrderedLocusNames=YALIOD08734g;
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Eukaryota; Fungi; Ascomycota; Saccharomycotina;
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Q6C9S5;
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Pro; IPR013256; Chromatin_SPT2.
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                                                                                                                                                                                                                                                    0;
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                                                                                                                                                                                                                                                 Gaps
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RESULT
BAMT_AN
ID BA
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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                InterPro; IPR000169; Pept_cys_AS.
InterPro; IPR013128; Peptidase_CIA.
InterPro; IPR0131201; Peptidase_CIA.
InterPro; IPR013201; Propeptide I29.
InterPro; IPR013201; Propeptide I29.
PANTHER; PTHR12411; Peptidase_CI; 1.
Pfam; PP08246; Inhibitor_I29; 1.
Pfam; PP00112; Peptidase_CI; 1.
Pfam; PP00112; Peptidase_CI; 1.
PRINTS; PR00705; PAPAIN.
PRODOM; PD000158; Peptidase_CI; 1.
SMART; SM00645; Peptidase_CI; 1.
SMART; SM00645; Peptidase_CI; 1.
PROSITE; PS000649; THIOL_PROTEASE_ASN; 1.
PROSITE; PS000639; THIOL_PROTEASE_HIS; 1.
PROSITE; PS00639; THIOL_PROTEASE_CYS; 1.
PROSITE; PS00639; THIOL_PROTEASE_CYS; 1.
PROSITE; PS00639; THIOL_PROTEASE_CYS; 1.
                                                                                                                                                                                                                                                                                                        ANTMA
BAMT_ANTMA
                    NUCLEOTIDE
STAGE, AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACT_SITE
ACT_SITE
DISULFID
DISULFID
DISULFID
                                                                                                                                            Antirrhinum majus (Garden Eukaryota; Viridiplantae;
                                                                                                                                                                                                               07-FEB-2006, entry version 15.
Benzoate carboxyl methyltransferase
                                                                                                                                                                                                                                                                     Q9FYZ9;
16-AUG-2005,
                                                                                                                            Spermatophyta;
                                                                                                                                                                                 Name=BAMT;
                                                                                                                                                                                                methionine:benzoic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Aliergen;
SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Muta E., Aramaki H., Takata Y., Kono A., Okamoto Y., Ota "Cloning and sequencing of fruit bromelain." Submitted (JAN-1993) to the EMBL/GenBank/DDBJ databases.
-I- CATALYTIC ACTIVITY: Broad specificity for cleavage o but strong preference for Z-Arg-Arg-|-NHMec among sm
       MEDLINE=20312758;
                                                                                                             asterids;
                                                                                                                                                                                                                                                  01-MAR-2001, sequence version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CHAIN
                                                                      NCBI_TaxID=4151;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EMBL; D14059;
PIR; T10503;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN=cv. N67-10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PROPEP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEROPS; C01.028;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Copyrighted by the UniProt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ++
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ALLERGEN: Causes an allergic reaction in human. SIMILARITY: Belongs to the peptidase C1 family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          substrates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            P07711;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6
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                                                                                                           lamiids;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hydrolase;
SEQUENCE [MRNA], PARTIAL PROTEIN SEQUENCE, DEVELO TISSUE SPECIFICITY.
312758; PubMed=10852939; DOI=10.1105/tpc.12.6.949;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  122
                                                                                                                                                                                                                                                                       integrated into UniProtKB/Swiss-Prot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1CJL.
                                                                                                       Magnoliophyta; eudicotyledons; ids; Lamiales; Plantaginaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BAA21849.1; -; mRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A
A
                                                                                                                                                                                                                                                                                                          STANDARD;
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279
184
217
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121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    78.7%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Protease;
                                                                                                                                                                                              carboxyl
                                                                                                                    snapdragon).
Streptophyta; Embryo
yta; eudicotyledons;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ۲.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bromelain.
/FTId=PRO 0000045810
By similarity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pred. No. 7.8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 37; DB 1;
Pred. No. 7.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     왕
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             By similarity.
/FTId=PRO_0000045809
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Potential.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    y similarity.
y similarity.
33781C55144242BO CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SE_HIS; 1.
Signal; Thiol protease; Zymogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          similarity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              similarity.
                                                                                                                                                                                              methyltransferase).
                                   PROTEIN SEQUENCE, DEVELOPMENTAL
                                                                                                                                                                                                                                                                                                        364 AA.
                                                                                                       Embryophyta; Tracheophyta; sdons; core eudicotyledons; ceae; Antirrhineae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 351;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          small
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of proteins
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RESULT
Q53LK1
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-MAY-2005, integrated into UniProtKB/TrEMBL.
24-MAY-2005, sequence version 1.
27-FEB-2006, entry version 3.
P-box domain, putative.
Oryza sativa (japonica cultivar-group).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; BEP clade;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPBIAS
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Plant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Buell C., Yuan Q., Ouyang S., Liu J., Wang A., Maiti R., Lin Zhu W., Hamilton J., Jones K., Tallon L., Feldblyum T., Tsit: Bera J., Kim M., Jin S., Fadrosh D., Vuong H., Overton II L. Reardon M., Weaver B., Johri S., Lewis M., Utterback T., Van Wortman J., Haas B., Koo H., Zismann V., Hsiao J., Iobst S., de Vazeilles A., White O., Salzberg S., Fraser C.; Submitted (JUN-2003) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Dudareva N., Murfitt L.M., Mann C.J., Gorenstein N., Kolosova N., Kish C.M., Bonham C., Wood K.; Kish C.M., Bonham C., Wood K.; "Developmental regulation of methyl benzoate biosynthesis and emission in snapdragon flowers.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                InterPro; IPR005299; Methytransf_6.
Pfam; PF03492; Methyltransf_7; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Copyrighted by the UniProt Consortium, Distributed under the Creative Commons
                                  Pfam; PF00646; F-box; 1.
SMART; SM00256; FBOX; 1.
                                                                                                                                             EMBL; AC145321; AAX95043.1; -;
                                                                                                                                                                                                                     Distributed
                                                                                                                                                                                                                                                        Copyrighted
                                                                                                                                                                                                                                                                                                                                Submitted (APR-2005)
                                                                                                                                                                                                                                                                                                                                                                                                     NUCLEOTIDE SEQUENCE.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=39947;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ehrhartoideae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Q53LK1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Q53LK1_ORYSA
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                                                                                                          InterPro; IPR001810; F-box.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        84
ORYSA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PTM: The N-terminus is blocked. SIMILARITY: Belongs to the meth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    manner, attracts the pollinators.
TISSUE SPECIFICITY: Expressed only in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and ovaries.

DEVELOPMENTAL STAGE: Expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FUNCTION: Converts benzoic acid into the volatile ester benzoates. This scent, mostly produced in a rhytmical, c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  lobes. Not found in the corolla tubes,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             289
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364 AA;
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41011 MW;
                                                                                                                                                                                                                                                            UniProt
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                                                                                                                                                                                                                                                                                                                                    to the
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75.0%;
                                                                                                                                                                                                                     iProt Consortium, see http://www.uniprot.org/terms
Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the methyltransferase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ٠.
                                                                                                                                                                                                                                                                                                                                    EMBL/GenBank/DDBJ databases
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Poly-Asp.
C10C8E864A581419 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
                                                                                                                                             Genomic_DNA
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RESULT 85
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Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                  PubMed=14704707; DOI=10.1038/nbt923;
Larimer F.W., Chain P., Hauser L., Lamerdin J.E., Malfatti S., Do I Land M.L., Pelletier D.A., Beatty J.T., Lang A.S., Tabita F.R., Gibson J.L., Hanson T.E., Bobst C., Torres y Torres J.L., Peres C., Harrison F.H., Gibson J., Harwood C.S.;
"Complete genome sequence of the metabolically versatile photosynthetic bacterium Rhodopseudomonas palustris.";
Nat. Biotechnol. 22:55-61(2004).
-i-FUNCTION: Involved in protein export. Acts as a chaperone by maintaining the newly synthesized protein in an open conformations in the newly synthesized protein in an open conformation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TIG_RHOPA
Q6N5L2;
                                                                                                   DOMAIN
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-JUN-2005, integrated into UniProtKB/Swiss-Prot. 05-JUL-2004, sequence version 1. 07-MAR-2006, entry version 10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                          CHAIN
                                                                                                                                                                                                                  Pfam; PF00254; FKBP_C; 1.
Pfam; PF05698; Trigger_C; 1.
Pfam; PF05697; Trigger_N; 1.
                                                                                                                                                                                                                                              INAMAP; MF 00303; -; 1.

IAMAP; MF 00303; -; 1.

InterPro; IPR001179; FKBP PPlase.

InterPro; IPR005215; Tri9_fac.

InterPro; IPR008880; Tri9ger_C.

InterPro; IPR008881; Trigger_N.
                                                                                                                                                                                                                                                                                                                      BioCyc;
                                                                                                                                                                                                                                                                                                                                                                         Copyrighted by the UniProt Consortium, see http://www.uniprot.org/teDistributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAIN=CGA009
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Name=tig; OrderedLocusNames=RPA2962;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Trigger factor (TF)
                                                                                                                                                                                         PIRSF, PIRSF003095; Trigger factor; TIGRFAMs; TIGR00115; tig; 1.
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                                                                                                                                                                                                                                                                                                                                               EMBL; BX572602;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rhodopseudomonas palustris
                                                                                                                                                                  Cell cycle; Cell division;

    -!- SIMILARITY: Belongs to the FKBP-type PPIase fam
    -!- SIMILARITY: Contains 1 PPIase FKBP-type domain.

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 YKDEDDK 449
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BX571963_GR; RPA2962.
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/FTId=PRO_
                                                               Score 37; DB 1;
Pred. No. 1e+03;
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9DCE13323ABB5800 CRC64;
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RESULT 86
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AC Q2R94
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DT 24-JJ
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Lafontaine I., de Montigny J., Marck C., Neuveglise C., Talla E.,
A Goffard N., Frangeul L., Aigle M., Anthouard V., Babour A., Barbe V.,
A Barnay S., Blanchin S., Beckerich J.-M., Beyne E., Bleykasten C.,
A Boisrame A., Boyer J., Cattolico L., Confanioleri F., de Daruvar A.,
A Despons L., Fabre E., Fairhead C., Ferry-Dumazet H., Groppi A.,
A Hantraye F., Hennequin C., Jauniaux N., Joyet P., Kachouri R.,
A Kerrest A., Koszul R., Lemaire M., Lesur I., Ma L., Muller H.,
A Nicaud J.-M., Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,
A Pellenz S., Potier S., Richard G.-F., Straub M.-L., Suleau A.,
A Swennen D., Tekaia F., Wesolowski-Louvel M., Westhof E., Wirth B.,
A Zeniou-Meyer M., Zivanovic Y., Bolotin-Fukuhara M., Thierry A.,
A Wincker P., Souciet J.-L.,
A Wincker P., Souciet J.-L.,
A Wincher S., Caudron B., Scarpelli C., Gaillardin C., Weissenbach J.,
A Wincker P., Souciet J.-L.,
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75_DEBHA
Q6BV75_DEBHA
Q6BV75;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC STRAIN=ATCC 36239 / CBS 767;
PubMed=15229592; DOI=10.1038/nature02579;
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24-JAN-2006,
24-JAN-2006,
07-FEB-2006,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Debaryomyces hansenii (Yeast) (Torulaspora hansenii) Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetales; Saccharomycetaceae; Debaryomyces
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ORPNames=LOC Os11g09970;

Oryza sativa (japonica cultivar-group).

Eukaryota; Viridiplantae; Streptophyta; lispermatophyta; Magnoliophyta; Liliopsida

Enrhartoideae; Oryzaae; Oryza.
                                                          Copyrighted
Distributed
                                                                                                                                            Nature 430:35-44(2004).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OrderedLocusNames=DEHA0C05478g;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarities with CA0612
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-AUG-2004, integrated into UniProtKB/TrEMBL.
16-AUG-2004, sequence version 1.
07-FEB-2006, entry version 10.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Buell C.R., Wing R.A., McCombie W.A., Ouyang S.; Submitted (APR-2005) to the EMBL/GenBank/DDBJ databases.
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                                                    by the UniProt Consortium, see http://www.uniprot.org/terms under the Creative Commons Attribution-NoDerivs License
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85.7%;
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-; Genomic_DNA
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Liliopsida; Poales; Poaceae; BEP cla
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F65EC789BB4635F4 CRC64;
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RESULT
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                        12_HPV20

VE2_HPV20

STANDARD;

P50766;

01-OCT-1996, integrated into Un
01-OCT-1996, sequence version 1
07-FEB-2006, entry version 23.
                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
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10-MAY-2005, sequence
07-FEB-2006, entry ver
Hypothetical protein A
Name=At1948400;
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W8_ARATH
Q56XW8_ARATH
                  Human papillomavirus ty
Viruses, dsDNA viruses,
                                                                Name=E2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SMART; SM00579; FBD; 1.
SMART; SM00256; FBOX; 1.
PROSITE; PS00430; TONB_DEPENDENT_REC_1; UNKNOWN_1.
                                                                                   Regulatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      InterPro; IPR001810; F-box.
InterPro; IPR006566; FBD.
InterPro; IPR013101; LLR 2.
InterPro; IPR010916; TONB Box N.
Pfam; PF00646; F-box; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Totoki Y., Seki M., Ishida J., Nakajima M., Enju A., Kami Narusaka M., Shin-i T., Nakagawa M., Sakamoto N., Oishi K Kobayashi M., Toyoda A., Sakaki Y., Sakurai T., Iida K., Satou M., Toyoda T., Konagaya A., Carninci P., Kawai J., Hayashizaki Y., Shinozaki K., "Large-scale analysis of RIKEN Arabidopsis full-length (R Submitted (MAR-2005) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hypothetical SEQUENCE 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Distributed under the Creative Commons
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=3702;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Arabidopsis thaliana (Mouse-ear cress).

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicotyledons;
rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Q56XW8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBL; AK221555; BAD94953.1; -; mRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUCLEOTIDE SEQUENCE.
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487 AA; 5
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                                                                                                                                                      integrated into UniProtKB/Swiss-Prot
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                                                                                       version
E2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       55627 MW;
                                       type
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Pred. No. 1.1e-
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                               Score 37; DB 2;
Pred. No. 1.1e+03;
0; Mismatches 1
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                    stage;
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                    Papillomaviridae,
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1.1e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 487;
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Oishi K.,
                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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Kohara Y.,
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InterPro; IPR010427; E2 C.
InterPro; IPR001866; E2 N.
Pfam; PF00511; PPV E2 C; 1.
Pfam; PF00508; PPV E2 C; 1.
ProDom; PD00672; E2 C; 1.
ProDom; PD00672; E2 C; 1.
ProDom; PD000672; E3 C; 1.
ProDom; PD000672; E3 C; 1.
ProDom; PD000672; E3 C; 1.
ProDom; PD000672; E2 C; 1.
ProDom; PD000672; E2 C; 1.
ProDom; PD000672; E3 C; 1.
ProDom; PD000672; E3 C; 1.
ProDom; PD000672; E3 C; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Vysotskia V.S., Schwartz J.R., Yu G., Toriumi M., Lenz C., Li Lee J.M., Li J., Li J., Gonzale A., Liu A., Liu K., Vaysberg Sakano H., Chin C., Choi E., Chiou J., Altafi H., Araujo R., Brooks S., Buehler E., Chao Q., Conn L., Conway A.B., Dunn P., Hansen N., Howng B., Huizar L., Khan S., Kim C., Palm C., Row Shinn P., Walker M., Davis R.W., Ecker J.R., Federspiel N.A., Theologis A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-MAY 2000, sequence version 1.
07-FEB-2006, entry version 23.
F11A17.5 protein.
Name=F11A17.5; OrderedLocusNames=At1g48400;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EMBL;
                                                                     InterPro; IPR006566; FBD.
InterPro; IPR013101; LLR 2.
InterPro; IPR0101916; TONE Box_N.
Pfam; PF00646; F-box; 1.
SMART; SM00579; FBD; 1.
SMART; SM00576; FBOX; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Arabidopsis thaliana (Mouse-ear cress)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicotyledons;
rosids; eurosids II; Brassloales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                               Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE
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[1]
                     PROSITE; PS50181; FBOX; 1.
PROSITE; PS50430; TONB_DEPENDENT_REC_1; UNKNOWN
                                                                                                                                                                                                                                                         TAIR; At1g48400;
                                                                                                                                                                                                                                                                                 EMBL; AC007932; AAD49757.1; -; PIR; B96524; B96524.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (JUN-2000) to the EMBL/GenBank/DDBJ databases.
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     513 AA;
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     58618 MW;
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85.7%;
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Pred. No. 1.1e
1; Mismatches
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     81A562DBB3DDA0F8 CRC64;
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Q59MN9;
26-APR-2005,
                                                                                                                                                                                                                                                                                                                                                                                                                     QEIEGE PLAF7
QEIEGE;
01-MAR-2003,
STRAIN=3D7;
Harris B., Lennard N., Clark L., Line A., Barron A., Corton Harris B., Lennard N., Clark L., Line A., Barron A., Corton Berriman M., Pain A., Hall N., Atkin R., Chillingworth C., Berriman M., Barrel Ormond D., Sanders M., Hayes R., Hall S., Quail M., Barrel Ormond D., Sanders M., Hayes R., Hall S., Quail M., Barrel Ormond D., Sanders M., Hayes R., Hall S., Quail M., Barrel Ormond D., Sanders M., Hayes R., Hall S., Quail M., Barrel Ormond D., Sanders M., Hayes R., Hall S., Quail M., Barrel Ormond D., Sanders M., Hayes R., Hall S., Quail M., Barrel Ormond D., Sanders M., Hayes R., Hall S., Quail M., Barrel Ormond D., Sanders M., Hayes R., Hall S., Quail M., Barrel Ormond D., Sanders M., Hayes R., Hall S., Quail M., Barrel Ormond D., Sanders M., Hayes R., Hall S., Quail M., Barrel Ormond D., Sanders M., Hayes R., Hall S., Quail M., Barrel Ormond D., Sanders M., Hayes R., Hall S., Quail M., Barrel Ormond D., Sanders M., Hayes R., Hall S., Quail M., Barrel Ormond D., Sanders M., Hayes R., Hall S., Quail M., Barrel Ormond D., Sanders M., Hayes R., Hall S., Quail M., Barrel Ormond D., Sanders M., Hayes R., Hall S., Quail M., Barrel Ormond D., Sanders M., Hayes R., Hall S., Quail M., Barrel Ormond D., Sanders M., Hayes R., Hall S., Quail M., Barrel Ormond D., Sanders M., Hayes R., Hall S., Quail M., Barrel Ormond D., Sanders M., Hayes R., Hall S., Quail M., Barrel M., Hayes R., Hall S., Quail M., Barrel M., Hayes R., Hall S., Quail M., Hayes R., Hall M., Hayes R., Hall M., Hayes R., Hall M., Hayes R., Hall M., Hayes R., Hal
                                                                                                                                                                                                                Plasmodium falciparum
Eukaryota; Alveolata;
NCBI_TaxID=36329;
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                                                                                                                                                                                                                                                                                                                Hypothetical protein MAL13P1.75.
Name=MAL13P1.75;
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07-FEB-2006, entry version 6.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms
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piel N.A., Chibana H., Dungan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EAK91730.1; -; EAK91744.1; -;
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Pred. No. 1.2e+03;
0; Mismatches 1
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                                                                                                              Corton
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                                                                                Doggett J.,
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XS Sasaki T., Matsumoto T., Yamamoto K., Sakata K., Baba T., Katayose Y.
XA Mu J., Niimura Y., Cheng Z., Nagamura Y., Antonio B.A., Kanamori H.,
XA Hosokawa S., Masukawa M., Arikawa K., Chiden Y., Hayashi M.,
XA Hijishita S., Honda M., Ichikawa Y., Idonuma A., Iijima M., Ikeda M.,
XA Ikeno M., Ito S., Ito T., Ito Y., Ito Y., Iwabuchi A., Kamiya K.,
XA Ikeno M., Ito S., Ito T., Ito Y., Ito Y., Iwabuchi A., Kamiya K.,
XA Karasawa W., Katagiri S., Kikuta A., Kobayashi N., Kono I.,
XA Nagasaki H., Nakashima M., Nakama H., Mizubayashi T., Mukai Y.,
XA Namiki N., Negishi M., Ohta I., Ono N., Saji S., Sakai K., Shibata M.
XA Namiki N., Nakashima M., Nakama Y., Nakamichi Y., Nakamura M.,
XA Namiki N., Yamagata H., Yamane H., Yoshiki S., Yoshihara R., Yukawa K.,
XA Shimokawa T., Shomura A., Song J., Takazaki Y., Terasawa K., Tsuji K.
XA Shimokawa T., Shomura A., Song J., Takazaki Y., Terasawa K., Tsuji K.
XA Shimokawa T., Shomura A., Song J., Takazaki Y., Terasawa K., Tsuji K.
XA Shimokawa T., Shomura A., Song J., Takazaki Y., Terasawa K., Tsuji K.
XA Shimokawa T., Shomura A., Song J., Takazaki Y., Terasawa K., Tsuji K.
XA Shimokawa T., Shomura A., Song J., Takazaki Y., Terasawa K., Tsuji K.
XA Shimokawa T., Shomura A., Song J., Takazaki Y., Terasawa K., Tsuji K.
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XA Shimokawa T., Shomura A., Song J., Takazaki Y., Terasawa K., Tsuji K.
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XA Shimokawa T., Shomura A., Song J., Takazaki Y., Terasawa K., Tsuji K.
XA Shimokawa T., Shomura A., Song J., Takazaki Y., Terasawa K., Tsuji K.
XA Shimokawa T., Shomura A., Song J., Takazaki Y., Terasawa K., Tsuji K.
XA Shimokawa T., Shomura A., Song J., Takazaki Y., Terasawa K., Tsuji K.
XA Shimokawa T., Shomura M., Taka M., T
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                                                                                                                                                                                                                                                                                                                                                                                    InterPro; IPR000005; HTHAraC.
InterPro; IPR004353; Yeast73DUF.
PANTHER; PTHR13027; Yeast73DUF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; AP003627; BAD87849.1; -; Genomic_DNA. EMBL; AP003228; BAD87032.1; -; Genomic_DNA. GO; GO:0006810; P:transport; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Spermatophyta; Magnoliophyta;
Ehrhartoideae; Oryzeae; Oryza,
NCBI_TaxID=39947;
                                                                                                                                                                                                                                                                                                                                                            Pfam; PF03164; DUF254;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oryza sativa (japonica cultivar-group).
Eukaryota; Viridiplantae; Streptophyta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Q5JL08_ORYSA PRELIMINARY;
Q5JL08;
15-FEB-2005, integrated into
15-FEB-2005, sequence version
07-FEB-2006, entry version 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; AL844509; CAD52291.1; -; Hypothetical protein. SEQUENCE 554 AA; 65670 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Copyrighted by the UniProt Consortium, see http://www.uniprot.org/toDistributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Name=P0459B04.31; Synonyms=P0020E09.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HSV-I stimulating-related protein-like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  93
ORYSA
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44
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                                                                                                                                                                                                                                                     PRO1546; YEAST73DUF.
; PSO0041; HTH_ARAC_FAMILY_1;
S 567 AA; 63751 MW; 1ABOBS
                                                                                                                              6;
                                                                                                                                                                Similarity
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                                                             DYKUDDDDK 8
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51
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75.0%;
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75
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Pred. No. 1.3e
1; Mismatches
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Pred. No. 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        eptophyta; Embryophyta; Tracheophyta;
Liliopsida; Poales; Poaceae; BEP cla
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                                                                                                                                                                                                                                                         1AB0B30FE1BE148B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 332A13B5174CDA85 CRC64;
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                                                                                                                                                                                                                                                                                           UNKCNOWN
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                                                                                                                                                      DB 2;
1.3e+03;
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                                                                                                                                                                                        Length 567;
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anamori H.,
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(., Tsuji
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M.-Y.,
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Best Local
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                                                                                                                                                                             Q5AAB5 CANAL PRELIMINARY;
Q5AAB5;
26-APR-2005, integrated into
26-APR-2005, sequence versio
07-FEB-2006, entry version 5
Hypothetical protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TIGREAMS;
PROSITE; F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Stein L.D., Bao Z., Blasiar D., Blumenthal T., Brent M.R., Chen N., Chinwalla A., Clarke L., Clee C., Coghlan A., Coulson A., D'Eustachio P., Fitch D.H.A., Fulton L.A., Fulton R.E., Griffiths-Jones S., Harris T.W., Hillier L.W., Kamath R., Kuwabara P.E., Mardis E.R., Marra M.A., Miner T.L., Minx P., Kuwabara P.E., Plumb R.W., Rogers J., Schein J.E., Sohrmann M., Spieth J., Stajich J.E., Wei C., Willey D., Wilson R.K., Durbin R., Waterston R.H.;
                                                           Eukaryota; Fungi; Ascomycota; Saccharomycetales; mitosporic NCBI_TaxID=237561;
                                                                                                                                    Candida albicans SC5314.
                                                                                                                                                            Hypothetical protein. ORFNames=CaO19.9993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Complete proteome; Hypothetical NON_TER 664 664
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; CAAC01000031; CAE62879, 1; -; Ge InterPro; IPR001678; Fmu NOLI/Nop2p. InterPro; IPR011023; Nop2p. InterPro; IPR011023; Nop2p. InterPro; IPR006174; rRNAsub_mtfrase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Copyrighted by the UniProt Consortium, see http://www.uniprot.org/teDistributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PLoS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      comparative genomics.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Nematoda; Chromadorea
Rhabditidae; Peloderinae; Caenorhabditis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Caenorhabditis briggsae.

Caenorhabditis briggsae.

Nematoda;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-NOV-2004,
07-FEB-2006,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Q61QD3_CAEBR
Q61QD3;
                       NUCLEOTIDE SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pfam; PF01189; Noll_Nop2_Fmu; 1.
ProDom; PD005242; NusB_region; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PubMed=14624247;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN=AF16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=6238;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             'The genome sequence of Caenorhabditis briggsae: a platform
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     94
CAEBR
                                                                                                                                                                                                                                                                                                                      CANAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           preliminary data.
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                                                                                                                                                                                                                                                                                                                                                                                                                94
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PS01153; NOL1_NOP2_
                                                                                                                                                                                                                                                                                                                                                                                                                DYSDDDD 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                664 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE [LARGE SCALE
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                                                                                                                                                                                                                                                                                              PRELIMINARY;
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in CBG07065
                       [LARGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             73746 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           78.7%;
85.7%;
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                       SCALE GENOMIC DNA
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Pred. No. 1
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                                                                                       Saccharomycotina; Saccharomycetes; Saccharomycetales; Candida.
                                                                                                                                                                                                                                                                                                 PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chromadorea;
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ome shotgun
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Genomic_DNA.
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Jones T., Federspiel N.A., Chibana H., Dungan J.,
Magee B.B., Newport G., Thorstenson Y.R., Agabian
Davis R.W., Scherer S.;
Q5AA27_CANAL
                                                                                                                                                                                                                                                                                                                 Tulman E.R., Afonso C.L., Lu Z., 'The genome of canarypox virus.";
J. Virol. 78:353-366(2004).
                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-JUL-2004, integrated into Un
05-JUL-2004, sequence version 1
07-FEB-2006, entry version 10.
CNPV322 ankyrin repeat protein.
                                                                                                                                                                                                       PRINTS; PR01415; ANKYRIN. SMART; SM00248; ANK; 5.
                                                                                                                                                                                                                                                                              Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                NUCLEOTIDE SEQUENCE.
STRAIN=ATCC VR-111;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "The diploid genome sequence of Candida albicans.";

Proc. Natl. Acad. Sci. U.S.A. 101:7329-7334 (2004).

-i- CAUTION: The sequence shown here is derived from

EMBL/GenBank/DDBJ whole genome shotgun (WGS) en
                                                                                                                                                             SEQUENCE
                                                                                                                                                                       ANK repeat;
                                                                                                                                                                                PROSITE; PS50297; ANK_REP_REGION; PROSITE; PS50088; ANK_REPEAT; 3.
                                                                                                                                                                                                                               InterPro; IPR002110; ANK. Pfam; PF00023; Ank; 6.
                                                                                                                                                                                                                                                       EMBL; AY318871; AAR83668.1; -;
                                                                                                                                                                                                                                                                                                                                         PubMed=14671117; DOI=10.1128/JVI.78.1.353-366.2004; Tulman E.R., Afonso C.L., Lu Z., Zsak L., Kutish G.F.,
                                                                                                                                                                                                                                                                                                                                                                                                                         Viruses; dsDNA viruses,
                                                                                                                                                                                                                                                                                                                                                                                                                                      Canarypox virus (CNPV).
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Name=CNPV322
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26
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Pro; IPR013154; ADH_N.
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Pred. No. 1.6e+03;
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RX PubMed=15875012; DOI=10.1038/nature03481;
RA Eichinger L., Pachebat J.A., Gloeckner G., Rajandream M.A.,
RA Eichinger L., Pachebat J.A., Gloeckner R., Szafranski K., Xu Q.,
RA Sucgang R., Berriman M., Song J., Olsen R., Szafranski K., Xu Q.,
RA Gunggang R., Berriman M., Bong J., Olsen R., Szafranski K., Xu Q.,
RA Tunggal B., Kummerfeld S., Madera M., Konforcov B.A., Rivero F.,
RA Fankier A.T., Lehmann R., Hamlin N., Davies R., Gaudet P., Fey P.,
RA Bankier A.T., Lehmann R., Hamlin N., Davies R., Davis P.,
RA Farbrother K., Chen G., Saunders D., Sodergren E.J., Davis P.,
RA Kerhornou A., Nie X., Hall N., Anjard C., Hemphill L., Bason N.,
RA Karhornou A., Nie Y., Hall N., Anjard C., Hemphill L., Bason N.,
RA Farbrother P., Desany B., Just E., Morio T., Rost R., Churcher C.M.,
RA Coper J., Haydock S., van Driessche N., Cronin A., Goodhead I.,
RA Coper J., Haydock S., van Driessche N., Harper D., Lindsay R.,
RA Muzny D.M., Mourier T., Pain A., Lu M., Harper D., Lindsay R.,
RA Muzny D.M., Mourier T., Pain A., Lu M., Harper D., Lindsay R.,
RA Hauser H., James K.D., Quiles M., Madan Babu M., Saito T.,
RA Mnights A., Loulseged H., Mungall K.L., Oliver K., Price C.,
RA Mights A., Loulseged H., Mungall K.L., Simmonds M.N., Spiegler S.,
RA Tivey A., Sugano S., White B., Walker D., Woodward J.R., Winckler T.,
RA Tanaka Y., Shaulsky G., Schleicher M., Weinstock G.M., Rosenthal A.,
RA Cox E.C., Chisholm R.L., Gibbs R.A., Loomis W.F., Platzer M.,
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Q54D84;
24-MAY-2005,
24-MAY-2005,
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26-APR-2005,
07-FEB-2006,
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Eukaryota; Fungi; Ascomycota;
Saccharomycetales; mitosporic
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Jones T., Federspiel N.A., Chibana H., Dungan J.,
Magee B.B., Newport G., Thorstenson Y.R., Agabian
Davis R.W., Scherer S.;
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ORFNames=CaO19.2457;
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InterPro; IPRO00210; BTB.
Pfam; PF00651; BTB; 1.
SMART; SM00225; BTB; 1.
PROSITE; PS50097; BTB; 1.
Hypothetical protein.
SEQUENCE 697 AA; 79148 M
                                                                                                                                                                                                                                                       Hypothetical proteIn.
SEQUENCE 700 AA; 78182 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "The diploid genome sequence of Candida albicans.";
Proc. Natl. Acad. Sci. U.S.A. 101:7329-7334(2004).
-I- CAUTION: The sequence shown here is derived from EMBL/GenBank/DDBJ whole genome shotgun (WGS) en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Magee B.B., Newport G., Thorstenson Y.R., Davis R.W., Scherer S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes; Saccharomycetales; mitosporic Saccharomycetales; Candida. NCBI_TaxID=237561;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hypothetical protein.
ORFNames=CaO19.10812, CaO19.3302;
Candida albicans SC5314.
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                                                                                                                                                                                                                                                                                                                      InterPro; IPR005036; CBM_21.
Pfam; PF03370; CBM_21; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PubMed=15123810; DOI=10.1073/pnas.0401648101; Jones T., Federspiel N.A., Chibana H., Dungan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hypothetical
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-APR-2005,
07-FEB-2006,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CAUTION: The sequence shown here is EMBL/GenBank/DDBJ whole genome shoto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   preliminary data.
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168
                                                                                                                                                                                                                                                                                                                                                                                         AACQ01000045; EAK99193.1; -;
AACQ01000042; EAK99468.1; -;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       437
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                                                                                                                                                             78.7%;
85.7%;
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                                                                                                                          Score 37; DB
Pred. No. 1.66
0; Mismatches
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Pred. No. 1.6e+03;
1; Mismatches 1
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R., Agabian
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                                                                                                                                                                                            DB 2;
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                                                                                                                                                             6e+03;
                                                                                                                                                                                     Length 700;
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RESULT 100
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X PubMed=15994558; DOI=10.1126/science.1110439;

A Gardner M.J., Bishop R., Shah T., de Villiers E.P., Carlton J.M.,

A Gardner M.J., Bishop R., Shah T., de Villiers E.P., Carlton J.M.,

A Hall N., Ren O., Paulsen I.T., Pain A., Berriman M., Wilson R.J.,

RA Sato S., Ralph S.A., Mann D.J., Xiong Z., Shallom S.J., Weidman J.,

RA Jiang L., Lynn J., Weaver B., Shoaibi A., Domingo A.R., Wasawo D.,

RA Crabtree J., Wortman J.R., Haas B., Angiuoli S.V., Creasy T.H., Lu C.,

RA Suh B., Silva J.C., Utterback T.R., Feldblyum T.V., Pertea M.,

Suh B., Silva J.C., Utterback T.R., Feldblyum T.V., Pertea M.,

RA Allen J., Nierman W.C., Taracha E.L., Salzberg S.L., White O.R.,

RA Fitzhugh H.A., Morzaria S., Venter J.C., Fraser C.M., Nene V.,

"Genome Sequence of Theileria parva, a Bovine Pathogen That Transforms

RT Lymphocytes.",

RE Science 309:134-137(2005).
                                                                                                      Best Loc
Matches
                                                                                                                                                   Query Match
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SEQUENCE
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02-AUG-2005,
07-FEB-2006,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gardner M., Bishop R., Shah T., de Villiers E., Carlton J.M., Hall N. Ren Q., Paulsen I.T., Pain A., Berriman M., Wilson R.J.M., Sato S., Ralph S.A., Mann D.J., Xiong Z., Shallom S.J., Wedidman J., Jiang L., Lynn J., Weaver B., Shoaibi A., Wasawo D., Crabtree J., Wortman J.R., Haas B., Angluoli S., Creasy T.H., Lu C., Suh B., Silva J.C., Utterback T., Feldblyum T., Pertea M., Allen J., Taracha E.L., Salzberg S.L., White O., Fitzhugh H.A., Morzaria S., Venter J.C., Salzberg S.L., White O., Fitzhugh H.A., Morzaria S., Venter J.C., Stabmitted (JUN-2005) to the EMBL/GenBank/DDBJ databases -i- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which is
                                                                                                                                                                                                                                                    InterPro; IPR001410; DEAD. Pfam; PF07717; DUF1605; 1. Pfam; PF04408; HA2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                          EMBL; AAGK01000001; EAN33347.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms
Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN=Muguga;
Gardner M., Bishop R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RNA helicase, putative ORFNames=TP01_0103; Theileria parva.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=5875;
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                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                                  GO:0005524; F:ATP binding; IEA.
GO:0008026; F:ATP-dependent helic.
GO:0000166; F:nucleotide binding;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              preliminary data.
527
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                                                                                                   Similarity 6; Conserv
YKDEDDK
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6.
                                                                                                Score 37; DB 2;
Pred. No. 1.6e+03;
1; Mismatches 0
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| 47 100.0 26 1 47 100.0 29 5 47 100.0 30 1 47 100.0 34 2 | 47 100.0 23 2 47 100.0 24 2 47 100.0 26 1 | 47 100.0 22 2 47 100.0 23 2 | 47 100.0 22 2 47 100.0 22 2 | 47 100.0 22 2 | 47 100.0 22 2 | 47 100:0 22 1 | 47 100.0 21 2 | 47 100.0 21 2 | 47 100.0 21 2 | 47 100.0 18 1 | 47 100.0 17 2 | 47 100.0 17 2 | 47 100.0 17 1 | 47 100.0 16 2 | 47 100.0 16 2 | 47 100.0 15 2 | 47 100.0 14 2 | 47 100.0 14 2 | 47 100.0 14 2 | 47 100.0 14 2 | 47 100.0 14 1 | 47 100.0 14 1 | 47 100.0 14 1 | 47 100.0 14 1 | 47 100.0 13 7 | 47 100.0 13 5 | 47 100.0 13 2 | 47 100.0 13 2 | 47 100.0 13 2 | 47 100.0 13 2 | 47 100.0 13 1 | 47 100.0 13 1 | 47 100.0 L3 L | 47 100.0 I3 I | 47 100.0 12 2 | 47 100.0 12 2 | 47 100.0 12 2 | 47 100.0 12 2 | 47 100.0 12 2 | 47 100.0 12 2 | 47 100.0 12 1 | 47 100.0 12 1 | 47 100.0 12 1 | 47 100 0 12 1 | 47 100.0 II 2 | 47 100.0 Ht 2 | 47 100 0 11 2 | 47 100.0 11 2 | 47 100.0 11 2 | 47 100.0 11 2 | 47 100.0 11 2 | 47 100.0 II 2 | 4' 100.0 10 b | 47 100.0 10 5 | 47 100 0 10 5 | 47 100.0 10 2 | 47 100.0 10 2 | 47 100.0 10 2 |
| 47 100.0 26 1 47 100.0 29 5 47 100.0 30 1 47 100.0 34 2 | 47 100.0 26 1 US-08-634-060-4 47 100.0 26 1 US-08-634-060- | 47 100.0 22 2 US-09-667-365- 47 100.0 23 2 US-09-270-767-3 | 47 100.0 22 2 47 100.0 22 2 | 47 100.0 22 2 US-09-270-767-: | 47 100.0 22 2 US-09-455-061-047 100.0 23 3 US-09-668-183 | 47 100.0 22 1 US-08-701-124 | 47 100.0 21 2 US-09-809-517A | 47 100.0 21 2 US-09-297-269- | 47 100.0 21 2 | 47 100.0 18 1 | 47 100.0 17 2 | 47 100.0 17 2 | 47 100.0 17 1 | 47 100.0 16 2 | 47 100.0 16 2 | 47 100.0 15 2 | 47 100.0 14 2 | 47 100.0 14 2 | 47 100.0 14 2 | 47 100.0 14 2 | 47 100.0 14 1 | 47 100.0 14 1 | 47 100.0 14 1 | 47 100.0 14 1 | 47 100.0 13 7 | 47 100.0 13 5 | 47 100.0 13 2 | 47 100.0 13 2 | 47 100.0 13 2 | 47 100.0 13 2 | 47 100.0 13 1 | 47 100.0 13 1 | 47 100.0 L3 L | 47 100.0 I3 I | 47 100.0 12 2 | 47 100.0 12 2 | 47 100.0 12 2 | 47 100.0 12 2 | 47 100.0 12 2 | 47 100.0 12 2 | 47 100.0 12 1 | 47 100.0 12 1 | 47 100.0 12 1 | 47 100 0 12 1 | 47 100.0 II 2 | 47 100.0 Ht 4 | 47 100 0 11 2 | 47 100.0 11 2 | 47 100.0 11 2 | 47 100.0 11 2 | 47 100.0 11 2 | 47 100.0 II 2 | 4' 100.0 10 b | 47 100.0 10 5 | 47 100 0 10 5 | 47 100.0 10 2 | 47 100.0 10 2 | 47 100.0 10 2 |
| 47 100.0 26 1 US-08-700-846-9 47 100.0 29 5 PCT-US94-05150-37 47 100.0 30 1 US-08-634-060-39 47 100.0 34 2 US-09-896-915-8 | 47 100.0 24 2 US-09-26-462-4 47 100.0 24 2 US-09-266-462-4 47 100.0 26 1 US-08-634-060-26 | 47 100:0 22 2 US-09-267-365-1948 47 100:0 23 2 US-09-270-767-34566 | 47 100.0 22 2 US-09-270-767-50661 47 100.0 22 2 US-09-809-517A-33 | 47 100.0 22 2 US-09-270-767-35444 | 47 100.0 22 2 US-09-455-061-29 | 47 100.0 22 1 US-08-701-124-29 47 100.0 22 2 US-09-130-225-29 | 47 100.0 21 2 US-09-809-517A-30 | 47 100.0 21 2 US-09-297-269-21 | 47 100.0 21 2 US-09-266-462-1 | 47 100.0 18 1 US-08-670-175-8 | 47 100.0 17 2 US-09-832-464-22 | 47 100.0 17 2 US-09-184-826-22 | 47 100.0 17 1 US-08-651-818A-22 | 47 100.0 16 2 US-09-252-656B-57 | 47 100.0 16 2 US-08-817-145-19 | 47 100.0 15 2 US-09-896-915-2 | 47 100.0 14 2 US-09-937-126-4 | 47 100.0 14 2 US-09-381-244B-5 | 47 100.0 14 2 US-08-469-419-403 | 47 100.0 14 2 US-08-764-114-403 | 47 100.0 14 1 US-08-871-161-9 | 47 100.0 14 1 US-08-469-319A-403 | 47 100.0 14 1 US-08-231-205A-9 | 4/ 100.0 | 47 100.0 13 7 5395760-16 | 47 100.0 13 5 PCT-US95-04228-41 | 47 100.0 13 2 US-10-009-332-33 | 47 100.0 13 2 US-09-982-610-41 | 47 100.0 13 2 US-09-299-495F-48 | 47 100.0 13 2 US-08-446-648-41 | 47 100.0 13 1 US-08-871-161-7 | 47 100.0 13 1 US-08-690-011A-48 | 47 100.0 L3 L US-08-231-2063-7 | 47 100.0 L3 L US-08-222-616-41 | 47 100.0 12 2 US-10-318-142-14 | 47 100.0 12 2 US-09-969-192-23 | 47 100.0 12 2 US-09-622-439-14 | 47 100.0 12 2 US-09-455-061-23 | 47 100.0 12 2 US-09-130-225-23 | 47 100.0 12 2 US-08-792-832A-9 | 47 100.0 12 1 US-08-709-177-53 | 47 100.0 12 1 US-08-701-124-23 | 47 100 0 12 1 IIS+08-709-173-53 | 47 100 0 13 1 110-08-350-884-53 | 47 100.0 | 47 100.0 11 3 10 10 10 10 10 10 10 10 10 10 10 10 10 | 47 100 0 11 2 IIS-09-848-816-147 | 47 100.0 11 2 US-08-375-199B-16 S | 47 100.0 11 2 US-09-964-201A-23 Se | 47 100.0 11 2 US-09-374-964-1 Se | 47 100.0 11 2 US-08-720-565-16 | 47 100.0 11 2 US-09-026-276-23 | 4/ 100.0 10 5 FC1:-U595-12960-1 | 47 100 0 10 5 ECT (COUT-04) 1 - 20 | 47 100 0 10 5 PCT-IIS93-04717-26 | 47 100.0 10 2 US-09-545-894-30 S | 47 100.0 10 2 US-09-631-531-10 Se | 47 100.0 10 2 US-09-033-524B-18 Se |
| 47 100.0 26 1 47 100.0 29 5 47 100.0 30 1 47 100.0 34 2 | 47 100.0 24 2 US-09-2/U-76/-49/83 Sequence 47 100.0 24 2 US-09-266-462-4 Sequence 47 100.0 26 1 US-08-634-060-26 Sequence | 47 100.0 22 2 US-09-667-365-1948 Sequence 47 100.0 23 2 US-09-270-767-34566 Sequence | 47 100.0 22 2 US-09-270-767-50661 Sequence 47 100.0 22 2 US-09-809-517A-33 Sequence | 47 100.0 22 2 US-09-270-767-35444 Sequence | 47 100.0 22 2 US-09-455-061-29 Sequence | 47 100.0 22 1 US-08-701-124-29 Sequence | 47 100.0 21 2 US-09-809-517A-30 Sequence | 47 100.0 21 2 US-09-297-269-21 Sequence | 47 100.0 21 2 US-09-266-462-1 Sequence | 47 100.0 18 1 US-08-670-175-8 Sequence | 47 100.0 17 2 US-09-832-464-22 Sequence | 47 100.0 17 2 US-09-184-826-22 Sequence | 47 100.0 17 1 US-08-651-818A-22 Segmence | 47 100.0 16 2 US-09-295-9138 Segmence | 47 100.0 16 2 US-08-817-145-19 Sequence | 47 100.0 15 2 US-09-896-915-2 Sequence | 47 100.0 14 2 US-09-937-126-4 Sequence | 47 100.0 14 2 US-09-381-244B-5 Sequence | 47 100.0 14 2 US-08-469-419-403 Sequence | 47 100.0 14 2 US-08-764-114-403 Sequence | 47 100.0 14 1 US-08-871-161-9 Sequence | 47 100.0 14 1 US-08-469-319A-403 Sequence | 47 100.0 14 1 US-08-231-205A-9 Segmence | 47 100.0 14 1 US-08-411-/958-403 Sequence | 47 100.0 13 7 5395760-16 Patent No | 47 100.0 13 5 PCT-US95-04228-41 Sequence | 47 100.0 13 2 US-10-009-332-33 Sequence | 47 100.0 13 2 US-09-982-610-41 Sequence | 47 100.0 13 2 US-09-299-495F-48 Sequence | 47 100.0 13 2 US-08-446-648-41 Sequence | 47 100.0 13 1 US-08-871-161-7 Segmence | 47 100.0 13 1 US-08-690-011A-48 Sequence | 4/ 100.0 13 1 US-08-331-3083-7 Sequence | 4/ 100.0 13 1 US-08-222-616-41 Sequence | 47 100.0 12 2 US-10-318-142-14 Sequence | 47 100.0 12 2 US-09-969-192-23 Sequence | 47 100.0 12 2 US-09-622-439-14 Sequence | 47 100.0 12 2 US-09-455-061-23 Sequence | 47 100.0 12 2 US-09-130-225-23 Sequence | 47 100.0 12 2 US-08-792-832A-9 Sequence | 47 100.0 12 1 US-08-709-177-53 Sequence | 47 100.0 12 1 US-08-701-124-23 Semience | 47 100.0 14 1 05-08-300-803-33 Sequence | 47 100 0 13 1 16-00-303-030-10 sequence | 47 100 0 11 3 176-063-656-16 Segmence | 47 100 0 11 2 10 10 10 11 11 1 | 47 100 0 11 2 IIS-00-848-515-147 Seminore | 47 100.0 11 2 US-08-375-1998-16 Semience | 47 100.0 11 2 US-09-964-201A-23 Seguence | 47 100.0 11 2 US-09-374-964-1 Sequence | 47 100.0 11 2 US-08-720-565-16 Sequence | 47 100.0 11 2 US-09-026-276-23 Sequence | 4/ 100.0 10 5 FCI-0595-1250-1 Sequence | 47 100 10 F DT. 11005 13060 1 | 47 100.0 10 5 PCT-IIS93-04717-26 Semience | 47 100.0 10 2 US-09-545-894-30 Sequence | 47 100.0 10 2 US-09-631-531-10 Sequence | 47 100.0 10 2 US-09-033-524B-18 Sequence |

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US-09-855-547-9
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US-09-962-756-2110
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Patent No. 5359039
GENERAL INFORMATION:
APPLICANT: Smith, Craig
APPLICANT: Goodwin, Raymond
TITLE OF INVENTION: Isolated Poxvii
TITLE OF INVENTION: Necrosis Facto
                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1
US-08-089-458B-7
ADDRESSEE: Patricia Anne Perkins, Imm
STREET: 51 University Street
CITY: Seattle
STATE: Washington
COUNTRY: U.S.A.
ZIP: 98101
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERAPING SYSTEM: Apple Operating Sys
SOFTWARE: Microsoft Word, Version #5.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/089,458B
FILING DATE: 07/09/93
CIASSIFICATION: 1NFORMATION:
NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2608
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
ADDRESSEE: Patricia Anne Perkins, Immunex Corporation
STREET: 51 University Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-638-610-4
US-08-959-584-4
US-08-959-584-4
US-08-959-378-4
US-08-959-378-4
US-08-959-378-4
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US-09-116-594-4
US-09-116-594-4
US-09-117-134-4
US-09-117-974-4
US-09-117-974-7
US-09-117-973-4
US-09-473-551-11
US-09-473-551-11
US-09-473-551-13
US-08-959-689A-4
US-09-473-551-13
US-08-959-689A-4
US-09-11A-21
US-09-299-495F-21
US-08-959-689A-3
US-08-959-118-2
US-08-959-118-2
US-08-959-118-2
US-08-959-118-2
US-08-959-118-2
US-08-959-119-3
US-08-9518-842-5
US-08-411-7958-315
                                                                                                                                                                                                                                                                                                                                             Poxvirus A53R-Equivalent Tumor Factor Antagonists
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Best Local Similarity
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US-08-323-474-4
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                                                              Query Match
Best Local
                                                  Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Patent No.
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APPLICANT: Ziegle
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                               NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                  REGISTRATION NUMBER: 32,172
REFERENCE/DOCKET NUMBER: 2609
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 587-0430
                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/905,600
FILING DATE: 26-JUN-1992
                                                                                                                   MOLECULE TYPE:
                                                                                                                                                                                 SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                   FILING DATE: 26-JUN-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: peptide
                                                              Local Similarity
                                                                                                                                        STRANDEDNESS:
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CLASSIFICATION:
                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                                       TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET:
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TOPOLOGY: linear
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amino acid
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51 University Street
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                                                  Conservative
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                                                                                                                               linear
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Pred. No.
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Pred. No. 5e+05;
                                                Mismatches
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                                                              5e+05;
                                                                      DB 1; Length 8;
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RESULT 3

RESULT 4 US-08-086-335C-12 ; Sequence 12, Application US/08086335C

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                                                          Matches
                                                                                     Query Match
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                                                                                                                                                                                                                                  TELEX: 756822
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/966,775
FILING DATE: 27-OCT-1992
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 907,224
FILING DATE: 01-JUL-1992
PRIOR APPLICATION NUMBER: US 899,660
APPLICATION NUMBER: US 899,660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                              MOLECULE TYPE: peptide IMMEDIATE SOURCE: CLONE: FLAG peptide
                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 02-JUN-
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 15-JUN-1992 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Microsoft Wo CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
                                                        Local Similarity
nes 8; Conserv
                                                                                                                                                                                            STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
                                                                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                                                                                                        TELEFAX:
                                                                                                                                                                                                                                                                                                                                             NAME: Seese, Kathryn A. REGISTRATION NUMBER: 32,172
                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                          ENGTH:
                              1 DYKDDDDK 8
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                                                                                                                                                                                                          amino acid
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                                                                                                                                                                                                                          8 amino acids
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Goodwin, Rayword
Smith, Craig A.
Armitage, Richard J.
Gruss, Hans-Jurgen
Gruss, Hans-Jurgen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        E: Kathryn A. Seese,
51 University Street
                                                          Conservative
                                                                                                                                                                                                                                                                                      (206) 233-0644
                                                                                                                                                                               linear
                                                                                                                                                                                                                                                                                                                                                                                              26-MAY-1992
                                                                                                                                                            peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                           02-JUN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12 APRIL 1994
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                                                                        100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Word, Version 5.1a
                                                        0,
                                                                          Pred. No.
                                                                                     Score 47;
                                                          Mismatches
                                                                                       Length 8;
                                                        Indels
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                                                        Gaps
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RESULT 5
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                                                                                                                                                                                                                                                                                     Sequence 5, Application US/08265086 Patent No. 5576191 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple 7.1
SOFTWARE: Microsoft Word, Version 5.1a
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/086,335C
FILING DATE: 01-UIL-1993
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: 756822
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: peptide HYPOTHETICAL: NO ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 587-0430
TELEFAX: (206) 233-0644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Price,
TITLE OF INVENTION:
                                                                                                                                                                                                  APPLICANT: Gayle, Margit
APPLICANT: Slack, Jennifer
APPLICANT: Gruss, Hans Juergen
APPLICANT: Sims, John E.
APPLICANT: Dower, Steven K.
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IMMEDIATE SOURCE:
CLONE: FLAG peptide
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                                                                                                                                                                    TITLE OF INVENTION: No. 5576191el Cytokine That Binds ST2
                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Anderson, Kathryn A. REGISTRATION NUMBER: 32,172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 51 Uni
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                                                                               CITY: Seattle
STATE: Washington
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                                                                 COUNTRY:
                                                                                                                    STREET:
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                                                    98101
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                                                                                                                  51 University Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                    USA
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                                                                                                                                      Immunex Corporation
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US-08-265-086-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 100.0%; Score 47; DB 1; Best Local Similarity 100.0%; Pred. No. 5e+05; Matches 8; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (206) 233-064
TELEX: 756822
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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NAME: Seese, Katnry....

REGISTRATION NUMBER: 32,172

REFERENCE/DOCKET NUMBER: 2825

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION 187-0430
                                                                                                          APPLICATION NUMBER: US/08/236,918A
FILING DATE: 06-May-1994
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/060,843
FILING DATE: 07-May-1993
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
HYPOTHETICAL: NO
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CLONE: FLAG peptide
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ATTORNEY/AGENT INFORMATION:
NAME: Anderson, Kathryn A.
REGISTRATION NUMBER: 32,172
REFERENCE/DOCKET NUMBER: 2801-B
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 587-0430
                                                                                                                                                                                                                                                                                                                        ZIP: 98101
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Alderson, Mark R. APPLICANT: Goodwin, Raymond G. APPLICANT: Smith, Craig A.
                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Cytokine Designated 4-1BB Ligand NUMBER OF SEQUENCES: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                      STREET: J.
STREET: J.
TTTY: Seattle
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FILING DATE: June 17, 1994
CLASSIFICATION: 530
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: Apple 7.1
SOFTWARE: Microsoft Word, Version 5.1a
                                                                                                                                                                                                                                                               COMPUTER: Apple Power Macintosh
OPERATING SYSTEM: Apple 7.5.3
SOFTWARE: Microsoft Word, Version #6.0.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
STRANDEDNESS: si
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                                                                                                                                                                                                                                                                                                                                                                                                                                     E: Kathryn A. Anderson, Immunex Corporation51 University Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMA., Kathryn A., Kathryn A., 32,172
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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                                                                                                                                                                                                                       FILING DATE: 01-JUL-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 907,224
                                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATION SYSTEM: Apple 7.1
SOFTWARE: Microsoft Word, Version 5.1a
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/570,923
ATTORNEY/AGENT INFORMATION:
NAME: Seese, Kathryn A.
REGISTRATION NUMBER: 32,172
REFERENCE/DOCKET NUMBER: 2804-E
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
                                                                                                                                    APPLICATION NUMBER: US 892,459
FILING DATE: 02-JUN-1992
PRIOR APPLICATION DATA:
                                                                                                                                                                                     APPLICATION NUMBER: US 899,660 FILING DATE: 15-JUN-1992 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                        CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple 7
                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Armitage, Richard J.
APPLICANT: Gruss, Hans-Jurgen
TITLE OF INVENTION: No. 5677430el Cytokine That Binds CD30
NUMBER OF SEQUENCES: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IMMEDIATE SOURCE:
CLONE: FLAG peptide
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HYPOTHETICAL: N
ANTI-SENSE: NO
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APPLICANT: Smith, Craig A.
                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/225,989
FILING DATE: 12 APRIL 1994
APPLICATION NUMBER: US 07/966,775
FILING DATE: 27-OCT-1992
                                                                                                     APPLICATION NUMBER: FILING DATE: 26-MAY
                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/0 FILING DATE: 12-DEC-1995 CLASSIFICATION: 530
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CITY: Seattle
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linear
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                                                                                                     UMBER: US 889,717
26-MAY-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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US-08-446-908-6
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                                                     FILING DATE: US 07/55, 209
FILING DATE: 13-APR-1995
FILING DATE: 13-APR-1994
APPLICATION NUMBER: US 07/957,649
FILING DATE: 21-APR-1994
APPLICATION NUMBER: US 07/957,649
FILING DATE: 06-OCT-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/511,438
FILING DATE: 13-APR-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/255,209
FILING DATE: 07-OCT-1988
PRIOR APPLICATION NUMBER: US 07/255,209
FILING DATE: 06-OCT-1987
APPLICATION NUMBER: US 07/113,566
FILING DATE: 06-OCT-1987
ATTORNEY/ACENT INFORMATION:
NAME: Seese, Kathryn A.
REGISTRATION NUMBER: 32,172
REFERENCE/DOCKET NUMBER: 32,172
REFERENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent No. 5705149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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Best Local (
TELEFAX: (206) 233-0644
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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TOPOLLOGY: lineal
MOLECULE TYPE: peptide
IMMEDIATE SOURCE:
TYPOLE: FLAG peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Microsoft Word, Version 5.1a CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/446 and FILING DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple 7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Goodwin, Raymond G. APPLICANT: Lupton, Stephen D. APPLICANT: Mochizuki, Diane Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Interleuk: TITLE OF INVENTION: Therewith NUMBER OF SEQUENCES: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Immunex Corporation
STREET: 51 University Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 DYKDDDDK 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (206) 233-0644
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                                 Query Match
Best Local Similarity
Thes 8; Conserve
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                                                                                                               ; MOLECULE TYPE:
US-08-634-060-25
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Best Local Similarity
Matches 8; Conserv
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MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTION PC-DOS/MS-DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/634,060
FILING DATE:
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 08/303,162
APPLICATION NUMBER: 08/303,162
APPLICATION NUMBER: 08/303,162
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                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 710
TELEFOMMUNICATION INFORMATION:
TELEFHONE: (312) 616-5600
TELEFAX: (312) 616-5700
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Kilyk, John Jr.
REGISTRATION NUMBER: 30763
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Leydig, V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Wickham, Thomas J.

APPLICANT: Kovesdi, Imre
APPLICANT: Roelvink, Petrus W.

TITLE OF INVENTION: ADENOVIRAL-MEDIATED CELL TARGETING COMMANDED
TITLE OF INVENTION: THE ADENOVIRUS PENTON BASE PROTEIN
NUMBER OF SEQUENCES: 60
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CLONE: FLAG peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: peptide HYPOTHETICAL: NO ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE: I
                                                                                                                                         LENGTH: 8 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET:
                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 08-SEP-1994
CLASSIFICATION: 514
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25, Application US/08634060
o. 5712136
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1 DYKDDDDK 8
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: Illinois
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Two Prudential Plaza, Suite 4900
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                                                 Conservative
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                                                               100.0%; Score 47; 100.0%; Pred. No.
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Pred. No. 5e+05;
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                                                 Mismatches
                                                               5e+05;
                                                                              DB 1; Length 8;
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RESULT 10
US-08-231-205A-6
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                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: US

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple MacIntosh
OPERATING SYSTEM: Apple 7.1
SOFTWARE: MICTOSOft Word, Version 5.1a
CURRENT APPLICATION DATA:
APPLICATION UNUMBER: US/08/231,205A
FILING DATE: 21-APR-1994
CLASSIFICATION UNMBER: US 07/957,649
FRIOR APPLICATION NUMBER: US 07/957,649
FRIOR APPLICATION NUMBER: US 07/511,438
APPLICATION NUMBER: US 07/511,438
APPLICATION NUMBER: US 07/511,438
FILING DATE: 13-APR-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/255,209
FRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/13,566
FILING DATE: 05-OCT-1988
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/113,566
FILING DATE: 26-OCT-1987
ATTORNEY/AGENT INFORMATION:
NAME: Seese, Kathryn A.
REGISTRATION NUMBER: 2104-D
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 587-0430
TELEPHONE: (206) 587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 6, Application US/08231205A Patent No. 5714585
                                                                            Matches
                                                                                              Best Local
                                                                                                               Query Match
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                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (206) 233-06
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                TOPOLOGY: lin
MOLECULE TYPE:
HYPOTHETICAL: N
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Intersewith
TITLE OF INVENTION: Therewith
THE OF SECULENCES: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Namen, Anthony E. APPLICANT: Goodwin, Raymond G. APPLICANT: Lupton, Stephen D. APPLICANT: Mochizuki, Diane Y.
                                                                                                                                                                       ANTI-SENSE: NO
IMMEDIATE SOURCE:
CLONE: FLAG peptide
                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS
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                                      1 DYKDDDDK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seattle
                                                                                              Similarity
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51 University Street
                                                                            Conservative
                                                                                                                                                                                                                                                                          linear
                                                                                                                                                                                                                                    NO
                                                                                                                                                                                                                                                  peptide
                                                                                                                                                                                                                                                                                    single
                                                                            100.0%; Score 47; DE 100.0%; Pred. No. 5et tive 0; Mismatches
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Interleukin-7 and Antibodies Reactive
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                                                                                                5e+05;
                                                                                                               DB 1; Length 8;
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                                                                                Gaps
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                             Sequence 3, Applic Patent No. 5726286
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                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: ALDERS
APPLICANT: ARMITA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: 1'

CTERISTICS:

B amino acid

STRANDEDNESS: a'

TOPOLOGY: 1'

TOPOLOGY: 1'
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PRIOR APPLICATION DATA:
APPLICATION UMBER: USSN 08/107,353
FILING DATE: 08-13-93
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Microsoft Word for Apple, Version : CURRENT APPLICATION DATA:
APPLICATION NUMBERS
                                                  APPLICANT: SPRIGGS, MELANIE
TITLE OF INVENTION: Isolated Epstein-Barr Virus BZLF2 Proteins
TITLE OF INVENTION: That Bind MHC Class II Beta Chains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Perkins, Patricia A
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 100
TELECOMMUNICATION INFORMATION:
             CORRESPONDENCE ADDRESS:
ADDRESSEE: IMMUNEX CORPORATION
                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Spriggs, Melanie
APPLICANT: Srinivasan, Subhashini
TITLE OF INVENTION: Methods of Preparing Soluble, Oligomeric
TITLE OF INVENTION: Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk COMPUTER: Apple Macintosh OPERATING SYSTEM: Apple OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET:
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No. 5716805
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                                                                                                COHEN, JEFFREY
COMEAU, MICHAEL
FARRAH, THERESA
SPRIGGS, MELANIE
SPRIGGS, MELANIE
51 UNIVERSITY STREET
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51 University Street
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ARMITAGE, RICHARD
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US-08-580-014-15
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INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: pepting IMMEDIATE SOURCE:
FILIMS .... 530
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/225,989
APPLICATION NUMBER: US/08/275,989
FILING DATE: 12 APRIL 1994
FILING DATE: US/07/966,775
                                                                                                    OPERATING SYSTEM: Apple 7.1
SOFTWARE: Microsoft Word, Version 5.1a
CURRINT APPLICATION DATA:
APPLICATION NUMBER: US/08/580,014
                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple 7
                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,693
REFERENCE DOCKET NUMBER: 26:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 08/2
FILING DATE: 04/28/94
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/430,633
FILING DATE: 28-APR-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity hes 8; Conserv
                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                            STREET:
                                                                                         FILING DATE:
                                                                                                                                                                                                                                                  STATE:
                                                                                                                                                                                                                                                                CITY:
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                                                                                                                                                                                                                                                                          E: Kathryn A.
51 University
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                                                                                                                                                                                                                                                                                                                                                                                     Goodwin, Raymond G.
Smith, Craig A.
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                                                                                       20-DEC-1995
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Pred. No. 5e+05;
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PRIOR APPLICATION DATA:

FILING DATE:

27-OCT-1992

APPLICATION NUMBER:

01-JUL-1992

US 907,224

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US-08-670-354-7
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APPLICANT: Steven R. Wiley and
APPLICANT: Raymond G. Goodwin.
TITLE OF INVENTION: Cytokine That Induces Apoptosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STKANNET INDEAL TOPOLOGY: lineal MOLECULE TYPE: peptide IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: 756822
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple 7.5.2
SOFTWARE: Microsoft Word, Version 6.0.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 26-MAY-1992
ATTORNEY/AGENT INFORMATION:
NAME: Seese, Kathryn A.
REGISTRATION NUMBER: 32,172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PILING DATE: 15-UN-1992
PRIOR APPLICATION DATA:
PROPRICATION NUMBER: US 892,459
                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 29-JUN-
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APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 9
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Local Similarity 100.0%;
hes 8; Conservative 0,
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                                                                                                                                                                                                                                                                                        STREET: 51 Un:
CITY: Seattle
                                                                                             APPLICATION NUMBER: US/08/670,354 FILING DATE: 25-JUN-1996 CLASSIFICATION: 435
                                                                                                                                                                                                                                                           COUNTRY:
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STRANDEDNESS: si
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APPLICATION NUMBER:
                              CLASSIFICATION: 435
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51 University Street
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                                          UMBER: US 08/496,632
29-JUN-1995
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US 08/548,368
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Pred. No.
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GENERAL INFORMATION:
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                                                                                                                                                                                              SOPTWARE: PATENTIN Release #1.24
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/442,043A
FILING DATE: 16-MAY-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/534,193
APPLICATION NUMBER: US 07/534,193
              FILING DATE: 05-JUN-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/9
FILING DATE: 24-AUG-1990
PRIOR APPLICATION NUMBER: US 07/9
PRIOR APPLICATION NUMBER: US 07/9
FILING DATE: 13-DEC-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/9
PRIOR APPLICATION NUMBER: US 07/9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 5.
STREET: 5.
CITY: Seattle
CTTATE: WA
TISA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Sims, John E.
APPLICANT: Cosman, David J.
APPLICANT: Lupton, Stephen D.
APPLICANT: Mosley, Bruce A.
APPLICANT: Mosley, Bruce A.
APPLICANT: Dower, Steven K.
TITLE OF INVENTION: Type II Interleukin-1 Receptors
NUMBER OF SEQUENCES: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HYPOTHETICAL: 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (200, 233-0644
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51 University Street
                                                                                                                                                                                                                                                                                                                                                                                                                                            USA
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DMBER: US 07/701,415
16-MAY-1991
                                                                                                                                          US 07/573,576
                                                                            US 07/627,071
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Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                            FILING DATE: 21-FEB-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Anderson, Kathryn A.
REGISTRATION NUMBER: 32,172
REFERENCE/DOCKET NUMBER: 2615
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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FILING DATE: 13-MAY-1994
ATTORNEY AGENT INFORMATION:
NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 20,693
REFERENCE/DOCKET NUMBER: 2003
TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-887-5570
                                                                                                         TELEX: 756822
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
TYPE: amino acid
STRANDENESS: sinc
TOPOLOGY: linear
MOLECULE TYPE: pept:
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 206-233-0644
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
FILING DATE: 21-FEB-19:
                                                                                                                                                                                                                                                                                                                                                                         COMPUTER REALABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Parnet, Patricia et al. TITLE OF INVENTION: Receptor Desig NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IMMEDIATE SOURCE:
CLONE: FLAG(r) peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
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FILING DATE: 12-JULY-1993
PRIOR APPLICATION DATA:
                                                                                                                                              TELEPHONE: (200) 233-0644
                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: Apple 7.1
SOFTWARE: Microsoft Word, Version 5.1a
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                                                                                          LENGTH:
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                                                                                          8 amino acids
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                                 ess: single
linear
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 . peptide
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                                                      Matches
                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 7, Application US/08308881
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APPLICANT: MOSLEY,
APPLICANT: COSMAN,
TITLE OF INVENTION:
                                                                                                                  MOLECULE TYPE: peptide
HYPOTHETICAL: NO
ANTI-SENSE: NO
IMMEDIATE SOURCE:
CLONE: FLAG peptide
                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Seese, Kathryn A.
REGISTRATION NUMBER: 32,172
REFERENCE/DOCKET NUMBER: 2614
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 587-0430
                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple 7.1
SOFTWARE: Microsoft Word, Version 5.1a
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/30B,881
FILING DATE: 12-SEP-1994
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 98101
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IMMEDIATE SOURCE:
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APPLICATION NUMBER: US 08/249,553
FILING DATE: 26-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: J.
CITY: Seattle
Thre: WA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                 TELEFAX: 12-
TELEFAX: 756822
                                                                                                                                                                                                                                                                               TELEPHONE: (205) 233-0644
                                                                   Local Similarity
                                                                                                                                                                                          STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                        TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLONE:
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1 DYKDDDDK 8
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                                                                  100.0%;
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                                                                Score 47; DB 1;
Pred. No. 5e+05;
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                                                                              DB 1; Length 8;
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RESULT 18 US-08-660-626-1

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                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Valeta Gregg
REGISTRATION NUMBER: 35,127
REFERENCE/DOCKET NUMBER: 0753
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 322-5070
TELEPAX: (415) 854-0875
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                               Sequence 1, Approximately No. 5801227
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Best Local :
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                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Fanslow, William C. III
APPLICANT: Zappone, Jodee
APPLICANT: Alderson, Mark
APPLICANT: Alderson, Mark
APPLICANT: Armitage, Richard J.
TITLE OF INVENTION: ANTIBODIES TO CD40
NUMBER OF SEQUENCES: 1
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCIII

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/660,626
PILING DATE:
CLASSIFICATION: 435
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APPLICANT: Stanley B. Prusiner
APPLICANT: Glenn C. Telling
APPLICANT: Fred E. Cohen
APPLICANT: Michael R. Scott
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple Operating System 7.1
SOFTWARE: Microsoft Word, Version 5.1a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
TYPE: amino acid
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ADDRESSEE: Fish & Richardson
STREET: 2200 Sand Hill Road, Suite 100
CITY: Menlo Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: TRANSGENIC ANIMALS EXPRESSING TITLE OF INVENTION: EPITOPE-TAGGED PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
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                                                                                                                                            STREET: 51
STREET: 51
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Best Local Similarity
                              CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION UNMER: 60/005,092
APPLICATION UNMER: 60/005,092
FILING DATE: 11-OCT-1995
ATTORNEY/AGENT INFORMATION:
NAME: GATES, EDWARD R,
REGISTRATION NUMBER: 31,616
REFERENCE/DOCKET NUMBER: B0801/7
TELEPHONE: 617-720-3500
TELEPHONE: 617-720-3441
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (206)233-064
INFORMATION FOR SEQ ID NO:
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APPLICATION NUMBER: US/08/130,541
FILING DATE: October 1, 1993
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                          ZIF: 02210
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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REGISTRATION NUMBER: 34.633
REFERENCE/DOCKST NUMBER: 28:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: WOLF, GREENFIELD & SACKS, P.C
STREET: 600 ATLANTIC AVENUE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: A NOVEL HAEMATOPOIETIC CYTOKINE AND USES TITLE OF INVENTION: THEREFOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: KIEFF, APPLICANT: DEVERG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ANTI-SENSE: NO
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SEQUENCE CHARACTERISTICS:
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FILING DATE:
CLASSIFICATION: 424
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TELEFAX: (206)233-0644
                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
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nilarity 100.0%;
Conservative 0
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BER: 2816
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Pred. No. 5e+05;
0; Mismatches 0;
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                                                        US-08-649-341A-9
                                                                          RESULT 22
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          Sequence 9, Application US/08649341A Patent No. 5847099 GENERAL INFORMATION:
                                                                                                                                                                                Matches
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Best Local (
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Best Local :
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Patent No. 5846782
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                            FILING DATE: 21-AUG-1996
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Wickham,
APPLICANT: Roelvink,
APPLICANT: Kovesdi,
                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IIM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/701,124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:

NAME/KEY: Binding-site

LOCATION: 1.8

LOCATION: /la
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HYPOTHETICAL: I
ORIGINAL SOURCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: Two Prud
CITY: Chicago
STATE: Illinois
                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: si
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les 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: /label= FLAG_EPITOPE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                               ENGTH:
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                                                                                                                                          1 DYKDDDDK 8
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                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                               8 amino acids
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Two Prudential Plaza - 49th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 USA
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Lin, Lih-Ling
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                                                                                                                                                                         100.0%; Score 47; DB 1; Length 8; llarity 100.0%; Pred. No. 5e+05; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Imre
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Best Local (
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OPERATING SYSTEM: PC COMPATIBLE
SOFTWARE: PATENTIN Release #1.0, Ver
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/649,341A
FILING DATE:
CLASSIFFATOR
                                              COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Va
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/494,440B
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CITY: Cambridge
CTATE: Massachusetts
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
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LENGTH: 8 amino acids
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NAME: Brown, Scott A,
REGISTRATION NUMBER: 32,724
                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Lin, Lih-Ling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: peptide
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MEDIUM TYPE: Floppy disk
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TITLE OF INVENTION: NOVEL THE RECEPTOR DEATH DOMAIN LIGAND
TITLE OF INVENTION: PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 498-8224
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                                                                                                                                                                         ZIP:
                                                                                                                                                                                                          CITY: Cambridge
STATE: Massachu
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8; Conserv
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Schievella, Andrea
SCHIEVELLA THE RECEPTOR DEATH DOMAIN LIGAND
NVENTION: PROTEINS AND INHIBITORS OF LIGAND BINDING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      E: Genetics Institute,
87 CambridgePark Drive
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REGISTRATION NUMBER:

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US-08-729-345-4
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                                Query Match
Best Local Similarity
Watches 8; Conserve
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GENERAL INFORMATION:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 498-8224
TELEPHONE: (617) 876-5851
INFORMATION FOR SEQ ID NO: 13:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
ATTORNEY AGENT INFORMATION:
NAME: Clark, Paul T
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 0484
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-542-8906
                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
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FILING DATE: 16-OCT-1996
CLASSIFICATION: 800
PRIOR APPLICATION DATA:
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MEDIUM TYPE: Diskette
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ADDRESSEE: Fish & Richardson P.C.
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APPLICANT: Berger-Sweeney, Joanne
TITLE OF INVENTION: TRANSGENIC NON-HUMAN MAMMAL
TITLE OF INVENTION: EXPRESSING FLAG-APP-C100 FUSION PROTEIN
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TYPE: amino acid
TOPOLOGY: linear
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SOFTWARE: FastSEQ Version 2.0
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                                                                                                                                                                                                                                                                              TOPOLOGY: lir
                                                                                                                                                                                                                                                                                                                                          TYPE: amino acids
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       1 DYKDDDDK 8
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                                                                       Conservative
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                                                                                                       100.0%;
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                                                                                                       Score 47;
Pred. No.
                                                                       Mismatches
                                                                                                5e+05;
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                                                                                                                                       DB 1;
                                                                                                                                   Length 8;
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RESULT 25
US-08-533-901B-15
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US-08-670-175-1
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APPLICANT: Lin, Lin-Ling
APPLICANT: Chen, Jennifer H.
APPLICANT: Schievella, Andrea
APPLICANT: Graham, James
APPLICANT: Graham, James
TITLE OF INVENTION: PROTEINS AN
                                                                                                                             GENERAL INFORMATION:
APPLICANT: LINDEN,
APPLICANT: TAYLOR,
APPLICANT: ROBEVA,
APPLICANT: WOODARD
APPLICANT: JIN, XI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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Patent No. 5854081
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Brown, Scott A,
REGISTRATION NUMBER: 32,724
REFERENCE/DOCKET NUMBER: G152
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 498-8224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (617) 876-58 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/533,901B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
             NUMBER OF SEQUENCES: 8

CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, ADDRESSEE: P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                          APPLICANT: JIN, XIAOWEI
TITLE OF INVENTION: STABLE EXPRESSION OF HUMAN ADENOSINE
TITLE OF INVENTION: RECEPTORS, AND ASSAYS EMPLOYING THE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15, Application US/08533901B
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GY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  E: Genetics Institute,
87 CambridgePark Drive
1755 S. JEFFERSON DAVIS HIGHWAY,
                                                                                                                                                LINDEN, JOEL
TAYLOR, HEIDI
ROBEVA, ANNA
WOODARD, ROBIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 USA
                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                WOODARD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (617) 876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Score 47; DB 1; Length 8; 100.0%; Pred. No. 5e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NOVEL THE RECEPTOR DEATH DOMAIN LIGAND PROTEINS AND INHIBITORS OF LIGAND BINDING
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Query Match
Best Local Similarity
""" hes 8; Conserv:
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US-08-777-405A-15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 703-413-2220 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
INFORMATION FOR SEQ ID NO:
                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/777,405A
FILING DATE:
CLASSIFICATION: 435
CTARSIFICATION: 435
                              ATTORNEY/AGENT INFORMATION:
NAME: NO. 558753and, Greta E.
REGISTRATION NUMBER: 35,302
REFERENCE/DOCKET NUMBER: 27866/33441
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 474-6300
TELEPAX: (312) 474-0448
                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Hoekstra, Merl F.
APPLICANT: Holtzman, Douglas A
TITLE OF INVENTION: No. 5858753el Lipid Kinase
NUMBBER OF SEQUENCES: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 24,618
REFERENCE/DOCKET NUMBER: 49
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION: NAME: OBLON, NORMAN F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                    STREET: 6300 S
CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY:
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FILING DATE: 20-JUN-1996
CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                  Illinois
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RESULT 28
US-08-620-694A-3
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Best Local S
Matches 8
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                                      Query Match
                                                                                                                                                                                                                                                                                                CLASSIFICATION: 435

PRIOR APPLICATION DATA:
APPLICATION UNBER: USSN 08/538,765
FILING DATE: 7 AUGUST 1995
CLASSIFICATION 1435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 08/410,535
FILING DATE: 23 MARCH 1995
CLASSIFICATION: 435
CLASSIFICATION: 435
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia Anne
REGISTRATION NUMBER: 34,695
REFERENCE/DOCKET NUMBER: 2617-B
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
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STRANDEDNESS: not
TOPOLOGY: not rele
MOLECULE TYPE: pept:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: Apple Power Macintosh
OPERATING SYSTEM: Apple Operating System 7.5.5
SOFTWARE: Microsoft Word for Apple, Version 6.0.1
CURRENT APPLICATION DATA:
                                                                                                            MOLECULE TYPE: p
                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Yao, Zhengbin
APPLICANT: Spriggs, Melanie
APPLICANT: Fanslow, William
TITLE OF INVENTION: No. 5869286el Receptor That Binds IL-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS
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STREET: 51 Univ
CITY: Seattle
y Match 100.0%; Score 47; DB 1; Length 8; Local Similarity 100.0%; Pred. No. 5e+05; hes 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                LENGTH: 8 amino acids
TYPE: amino acid
STRANDEDNESS: not rela
TOPOLOGY: linear
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                                                                                        CLONE: FLAG_ peptide
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Similarity 100.0%;
B; Conservative 0
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51 University Street
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                                                                                                                              peptide
                                                                                                                                                                    not relevant
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Pred. No. 5e+05;
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RESULT 29
US-08-720-258-7
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                                                                                           US-08-977-871A-15
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                                                                                                           RESULT 30
                                           Sequence 15, Application US/08977871A Patent No. 5882910 GENERAL INFORMATION:
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GENERAL INFORMATION:
                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                  Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US 08/5:
PRILING DATE: 20 DEC-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/5:
APPLICATION NUMBER: US 08/5:
PILING DATE: 29 SEP-1995
ATTORNEY/AGENT INFORMATION:
NAME: Anderson, Kathryn A.
REGISTRATION NUMBER: 32,172
                                                                                                                                                                                                                                                                          HYPOLILLE NU ANTI-SENSE: NU IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: |
HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: 32,172
REFERENCE/DOCKET NUMBER: 2620-B
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR TANTION NUMBER: US 08/575,715
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: Apple Power Macintosh
OPERATING SYSTEM: Apple System 7.5.3
SOFTWARE: Microsoft Word, Version 6.0.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Smith, CI
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
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                APPLICANT:
   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: FILING DATE: 26-SEI
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8; Conserv
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Chantry, David
Hoekstra, Merl F.
Holtzman, Douglas A
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RESULT 31
US-08-839-032A-19
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                                                                                                                                                                                                                                                                                                                                                           Sequence 19, Application US/08839032A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Lin, Lih-Ling
APPLICANT: Chen, Jennifer H.
APPLICANT: Schievella, Andrea
APPLICANT: Graham, James
TITLE OF INVENTION: NOVEL TNF
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NAME: No. 5882910and, Greta E
REGISTRATION NUMBER: 35,302
REFERENCE/DOCKET NUMBER: 2786
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 474-6300
                                         COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: PC-DOS/MS-DOS

OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: not relevant MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/977,871A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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             SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                             STREET: 8, __
CTTY: Cambridge
                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
ODERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (312) 474-0448
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 8 amore acid
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APPLICATION NUMBER:
                                                                                                               ZIP:
                                                                                                                             COUNTRY:
                                                                                                                                                                                              ADDRESSEE:
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E: Illinois
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6300 Sears Tower/233 South Wacker Drive
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US/08/839,032A
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Pred. No.
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                              Version #1.25
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Best Local
                                                                                                        TELEFAX: (206) 233-064
TELEX: 756822
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                ATTORNEY AGENT INFORMATION:
NAME: Seese, Kathryn A.
REGISTRATION NUMBER: 32,172
                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 587-0430
TELEPAX: (206) 233-0644
                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/308,881
FILING DATE: 12-SEP-1994
APPLICATION NUMBER: US 08/249,553
FILING DATE: 26-MAY-1994
               MOLECULE TYPE:
HYPOTHETICAL: 1
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Microsoft Word, Version 5.1a CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Mosley, Bruce
APPLICANT: Cosman, David J.
TITLE OF INVENTION: Receptor for Oncostatin
NUMBER OF SEQUENCES: 11
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TOPOLOGY: linear
MOLECULE TYPE: peptide
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LENGTH: 8 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS
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MMEDIATE SOURCE:
                                                           TYPE: amino acid
STRANDEDNESS: sin
TOPOLOGY: linear
                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
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No. 5891997
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8; Conserv
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51 University Street
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Pred. No. 5e+05;
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RESULT 34
US-08-936-854-3
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US-08-870-518-27
; Sequence 27, A
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            Sequence 3, Appli
Patent No. 592573
                                                                                                                                                                  Matches
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Best Local :
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Best Local :
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/870,5
FILING DATE: 06-JUN-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/019,219
FILING DATE: 06-JUN-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
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TELECOMMUNICATION INFORMATION:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Davis, Roger J.

APPLICANT: Galcheva-Gargova, Zoya
TITLE OF INVENTION: NON-ACTIVATED RECEPTOR COMPLEX
TITLE OF INVENTION: PROTEINS AND USES THEREOF
                                                                                                                                                                                                                                                                                                                       TELEFAX: 61...
TELEFAX: 200154
                                                                                                                                                                                                                                                         LENGTH: 8 amino acids
TYPE: amino acid
TOPOLOGY: linear
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OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Fasse, Peter J.
REGISTRATION NUMBER: 32,983
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8; Conserv
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                           Application US/08936854
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                                                                                                                                                              Score 47; DB 1;
Pred. No. 5e+05;
Mismatches
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Pred. No. 5e+05;
Mismatches
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APPLICANT:

ALDERSON, MARK ARMITAGE, RICHARD COHEN, JEFFREY

APPLICANT: SPRIGGS, MELANIE TITLE OF INVENTION: Isolate TITLE OF INVENTION: That Bi

Isolated Epstein-Barr Virus BZLF2 Proteins That Bind MHC Class II Beta Chains

APPLICANT:

COMEAU, MICHAEL FARRAH, THERESA

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Best Local S
Matches 8
                                                                                                                                                                                                                                                                           Sequence 7, Application US/09059099 Patent No. 5925740 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -08-936-854-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/430,633
FILING DATE: 28-APR-1995
APPLICATION NUMBER: 08/235,397
FILING DATE: 04/28/94
ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A.
REGISTRATION UNMBER: 34,693
REFERENCE/DOCKET NUMBER: 2612
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (206)233-0644
INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple Operating System 7.1
SOFTWARE: Microsoft Word for Apple, Version 5.1a
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/936,854
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: IMMUNEX CORPORATION
STREET: 51 UNIVERSITY STREET
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple 7.
                                                                                                                                                                                               APPLICANT: Mosley, Bruce
APPLICANT: Cosman, David J.
TITLE OF INVENTION: Receptor for Oncostatin
NUMBER OF SEQUENCES: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
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CLONE: FLAG_ peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: (206)587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: SEATTLE
                                                                             COUNTRY: (
                                                                                                                                                STREET:
                                                                                                                                                                 ADDRESSEE:
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8; Conserv
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                                                                                                                                                E: Immunex Corporation
51 University Street
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        peptide
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RESULT 36
US-08-713-928B-10
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                                                                                                                                                                                                                                                                                                                                                                                                                                  PATENT NO. 5929304

GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: RADIN, DAVID N.
APPLICANT: OISHI, KAREN K.
APPLICANT: WEISSENBORN, DEBORAH L.
APPLICANT: WEISSENBORN, DEBORAH L.
TITLE OF INVENTION: PRODUCTION OF LYSOSOMAL ENZYMES IN
TITLE OF INVENTION: PLANT-BASED EXPRESSION SYSTEMS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 10,
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Best Local Similarity
          COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC COMPUTER:
COMPUTER: IEM PC COMPUTER:
COMPUTER: PATENTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN DATA:
SOFTWARE: PATENTIN DATA:
APPLICATION NUMBER: US/08/713,928B
FILING DATE: 13-SEP-1996
CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION TOMBER: US/00/003,737
FILING DATE: 14-SEP-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEX: 756822
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
ANTI-SENSE: NO
IMMEDIATE SOURCE:
CLONE: FLAG peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PILING DATE: 26-MAY-1994
ATTORNEY AGENT INFORMATION:
NAME: Seese, Kathryn A.
REGISTRATION NUMBER: 32,172
REFERENCE/DOCKET NUMBER: 2614
TELECOMMUNICATION INFORMATION:
TELEPHONE: (296) 587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Microsoft Word, Version 5.1a CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/059,099
                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION DATA:
PRIOR APPLICATION NUMBER: US/08/308,881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (206) 233-0644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: | FILING DATE: 12-SEP-| APPLICATION NUMBER: |
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                 ZIP:
                                                                                                                                                                                                                                                                                                                       CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 8 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8;
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                                                                                                                                                                                                                                                                               10036-2711
Coruzzi, Laura A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Application US/08713928B
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                                                                                                                                                                                                                                                                                                      USA
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                                                                                                                                                                                Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-713-928B-10
                                                           Query Match
Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 1, Application US/08595043A Patent No. 5935824
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                 TELEPHONE: (415) 705-84:
TELEPAX: (415) 397-8338
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: SGARLATO, GREGORY D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/595,043A
                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: SG
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 705-8410
                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: CARROLL, PETER G.
REGISTRATION NUMBER: 32,837
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 90
CORRESPONDENCE ADDRESS:
ADDRESSEE: MEDLEN & CARROLL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: unknown MOLECULE TYPE: peptide
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9990
TELEPHAX: (212) 869-9741
                                                                                                                                              MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION:
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                                                                                                                                                                                STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 220 MONTGOMERY STREET, SUITE 2200
CITY: SAN FRANCISCO
STATE: CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                   TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 31-JAN
CLASSIFICATION: 435
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TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                 ENGTH:
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                                                                                                                                                                                                                   8 amino acids
                                                                 Conservative
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                                                                                                                                                                   unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Floppy disk
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                                                                             100.0%; Score 47; DB 1; 100.0%; Pred. No. 5e+05;
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                                                               Mismatches
                                                                                             Length 8;
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                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TOPOLOGY: linear; MOLECULE TYPE: peptide US-08-690-011A-50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-690-011A-50
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                                                                                                                                                                             Patent No. 5948638
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Lin, Lih-Ling
APPLICANT: Chen, Jennifer H.
APPLICANT: Schievella, Andrea
APPLICANT: Graham, James
TITLE OF INVENTION: NOVEL TNF RECEPTOR DEATH DOMAIN LIGAND
TITLE OF INVENTION: PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 15, Apr. Sequence 15,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 50 Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/690,011A
FILING DATE: 31-UUL-1996
PRIOR APPLICATION DATA:
APPLICATION UMBER: 60/001,654
FILING DATE: 31-UUL-1995
PRIOR APPLICATION UMBER: 60/018,496
FILING DATE: 29-MAY-1996
ATTORNEY/AGENT INFORMATION:
NAME: Serunian, Leslie A.
NAME: Serunian, Leslie A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (212)751-684
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: VINSON,
APPLICANT: KRYLOV,
                                                                  ADDRESSEE: Genetics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 35,353
REFERENCE/DOCKET NUMBER: 20
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30B
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 10154-0053
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS
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TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 8 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        50, 5942433
70. 5942433
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               h 100.0%; Score 47; DI Similarity 100.0%; Pred. No. 5e-8; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New York
Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Application US/08839031A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ϋ́
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Application US/08690011A
                               E: Genetics Institute,
87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 345 Park Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            : (212)758-4800
(212)751-6849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Floppy disk
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EXTENSION OF A PROTEIN-PROTEIN
INTERACTION SURFACE TO INACTIVATE THE FUNCTION OF A
CELLULAR PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Charles R.
                                                                                                                                                 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2026-4199US1
                                                                      Inc.
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STATE: Massachusetts COUNTRY: USA

02140

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US-08-762-106-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; TOPOLOGY: linear; MOLECULE TYPE: peptide US-08-839-031A-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 3, Application US/08762106 Patent No. 5948677
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Jarvik, Jonathan
TITLE OF INVENTION: READING
TITLE OF INVENTION: TAGGING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (617) 876-585
INFORMATION FOR SEQ ID NO:
                                                                                        TELEPHONE: (619) 654-2
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Sprunger, Suzanne A.,
REGISTRATION NUMBER: 41,323
REFERENCE/DOCKET NUMBER: GI5232BDIV
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 498-8284
TELEPAX: (617) 876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
COMPUTER: PATEN: PC-DOS/MS-DOS
SOFTMARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/839,031A
FILING DATE:
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UNBER: US/08/762,106
FILING DATE: 09-DEC-1996
CLASSIFICATION: 435
CTANSIFICATION: 435
                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Brotman, Harris F.
REGISTRATION NUMBER: 35,461
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 654-2428
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
MOLECULE TYPE:
FRAGMENT TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: La Jolla
STATE: California
                                   LENGTH: 8 amino a TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET:
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Similarity 100.0%;
8; Conservative 0
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                                                                        8 amino acids
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 internal
                   protein
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Pred. No. 5e+05;
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Query Match
Best Local Similarity
Theohes 8; Conserva
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US-09-016-366A-30
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RESULT 42
US-08-700-846-8
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Matches
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                                                                                                                                                                 Query Match
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APPLICANT: Stevens, Richard L.
APPLICANT: Huang, Chifu
TITLE OF INVENTION: MAST CELL PROTEASE PEPTIDE
TITLE OF INVENTION: INHIBITORS
                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION UMBER: 60/037,
FILING DATE: 05-FEB-1997
ATTORNBY/AGENT INFORMATION:
NAME: Plumer, Elizabeth R.
REGISTRATION UMBER: 36,637
REFERENCE/DOCKET NUMBER: B0
                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ for Windows Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/016,366A
FILING DATE: January 30, 1998
FILING DATE: January 30, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 02210-2211
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 65
CORRESPONDENCE ADDRESS:
ADDRESSEE: Wolf, Greenfield & Sacks, P.C
                                                                                                                                                                                                                TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: 617-720-3500
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STREET: VC
STREET: VC
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                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                  STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                TELEFAX: 617-720-2441
                                                                                                                                                                                                                                                                                   LENGTH:
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                                                                                                                               100.0%; Score 47; Di
ilarity 100.0%; Pred. No. 5e-
Conservative 0; Mismatches
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Sequence 8, Application US/08700846
Patent No. 5967311
GENERAL INFORMATION:
APPLICANT: WICKHAM, THOMAS J.
APPLICANT: ROELVINK, PETRUS W.

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RESULT 43
US-08-871-161-6
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                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Namen, Anthony E.
APPLICANT: Goodwin, Raymond G.
APPLICANT: Lupton, Stephen D.
APPLICANT: Mochizuki, Diane Y.
TITLE OF INVENTION: Interleukin-
TITLE OF INVENTION: Therewith
NUMBER OF SEQUENCES: 17
                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 6,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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FILING DATE: 21-AUG-1996
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: LARCHER, CAROL
REGISTRATION NUMBER: 35243
REFERENCE/DOCKET NUMBER: 7429.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 616-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO:
                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple 7.1
SOFTWARE: Microsoft Word, Version 5.1a
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
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CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 60601-6780
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: KOVESDI, TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: LEYDIG, V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
APPLICATION NUMBER: US/0 FILING DATE: 09-JUN-1997
                                                                                                                                                                                           COUNTRY: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                , Application US/08871161 5965122
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TWO PRUDENTIAL PLAZA, SUITE 4900
                                                                                                                                                             SD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        : (312) C16-5700 (312) 616-5700 8:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 47; DB 1; Length 8; ilarity 100.0%; Pred. No. 5e+05; Conservative 0; Mismatches 0; Indels
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Interleukin-7 and Antibodies Reactive
                US/08/871,161
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                                                                                                                                                                                                                                                                                                                                                                           Sequence 1, Application US/08832935 Patent No. 5965375 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 8; Conservat
                                                                                          ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, V.
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (206) 233-06.
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/
PILING DATE: 26-OCT-1987
PITORNEY/AGENT INFORMATION:
NAME: Seese, Kathryn A.
REGISTRATION NUMBER: 32,17
               ATTORNEY/AGENT INFORMATION: NAME: Smith, Timothy L.
                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                            APPLICANT: Valkirs, TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IMMEDIATE SOURCE:
CLONE: FLAG peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: 1i
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APPLICATION NUMBER: US 0:
FILING DATE: 07-OCT-1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ANTI-SENSE: NO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 06-OCT-PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 8 amino ac
TYPE: amino acid
STRANDEDNESS: sinc
                                              APPLICATION NUMBER: FILING DATE: 04-APF CLASSIFICATION: 435
                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                           CITY: San Francisco
STATE: California
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
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APPLICATION NUMBER:
FILING DATE: 21-API
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                                                                                                                                                                                                               USA
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                                                             04-APR-1997
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22-MAY-1995
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Pred. No. 5e+05;
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                                                                                                               Version #1.30
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                    Query Match
Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:

APPLICANT: Wang, Bruce
APPLICANT: Fisher, Joseph
APPLICANT: Payan, Donald
TITLE OF INVENTION: No. 5968802el Nuclear Cyclophilin
NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Patent No.
                                                                                                                                                                                                                                                                        TELEFAX: (415) 398-32
TELEX: 910 277299
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Silva, Robin M.
REGISTRATION NUMBER: 38,304
                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: A-61230/DJB/RMS
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
                                                                                                                                                               MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: C:
COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              y Match 100.0%; Score 47; DE
Local Similarity 100.0%; Pred. No. 5e+
hes 8; Conservative 0; Mismatches
                                                                                                                                                                              STRANDEDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/0 FILING DATE: 07-JUN-1995 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: Four Embarca
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Flehr, Ho
ADDRESSEE: & Herbert
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
                                                                                                                                                                                                                    TYPE: amino acid
                                                                                                                                                                                                                                     LENGTH:
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                                  рукоррок в
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                       8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            United States
                                                                                                                                                                                                                                                                                                              (415) 398-3249
                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                linear
                                                                                                                                                                                                   unknown
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                                                                    100.0%; Score 47; DB 1; Length 8; 100.0%; Pred. No. 5e+05; tive 0; Mismatches 0; Indels
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RESULT 46
US-08-876-882-5
                                                                                           Sequence 15, Application.

Sequence 15, Application.

Sequence 15, Application.

GENERAL INFORMATION:

APPLICANT: Hoekstra, Merl F.

APPLICANT: Holtzman, Douglas A.

APPLICANT: Holtzman, No. 5985589el Lipid Kinase

TITLE OF INVENTION: No. 5985589el Lipid Kinase
                                                                                                                                                                                                                                                            RESULT 47
US-09-225-951-15
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Patent No. 5981201
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: FASESEQ for Windows Version CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/876,882 FILING DATE: 16-JUN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 435
PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
ZIP: 02173-4799
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Groopman, TITLE OF INVENTION: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 36,361
REFERENCE/DOCKET NUMBER: NE
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 08-JAN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Doreen, Hogle M
REGISTRATION NUMBER: 36,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Avraham, Hava
APPLICANT: Groopman, Jero
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: IWC
STREET: Lexington
                                                              ADDRESSEE:
STREET: 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER:
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                         CITY: Chicago
STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE:
ZIP: 60606
                COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8 amino acids
                                                               E: Marshall O'Toole Gerstein Murray & Borun
6300 Sears Tower/233 South Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   781-861-9540
                USA
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METHODS OF DETECTION AND TREATMENT
OF BREAST CANCER
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                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 8;
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Query Match
Best Local Similarity
"hes 8; Conserv?
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 7, Application US/09058264 Patent No. 6010886
                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/308,881
FILING DATE: 12-SEP-1994
APPLICATION NUMBER: US 08/249,553
FILING DATE: 26-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: Seese, Kathryn A.
REGISTRATION NUMBER: 32,172
REFERENCE/DOCKET NUMBER: 2614-A
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 10F0RMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (312) 474-0441
TELEX: 25-3856
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:

NAME: NO. 5985589and, Greta E
REGISTRATION NUMBER: 35,302
REFERENCE/DOCKET NUMBER: 2786
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 474-6300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CUMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/225,951
FILING DATE:
FILING DATE:
                                                                                                                                                                                                                                                        OPERATING SYSTEM: Apple 7.1
SOFTWARE: Microsoft Word, Version 5.1a
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/058,264
                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple 7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Cosman, David J. TITLE OF INVENTION: Receptor for Oncostatin M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Mosley, Bruce
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 51 Uni
TELEPHONE: (200, 233-0644
                                                                                                                                                                                                                           FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (312) 474-0448
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
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51 University Street
                                                                                                                                                                                                                                                                                                                                                                                                       USA
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TYPE: amino acid;
; TOPOLOGY: linear;
; MOLECULE TYPE: peptide
US-08-991-426-8
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     Matches
               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 8, Application US/08991426
Patent No. 6013257
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
                                                                                                                                                                                                                                                                                             FILING DATE: 16-DEC-1997
PRIOR APPLICATION DATA: 08/851,160
APPLICATION UNMBER: 08/851,160
FILING DATE: 05-MAY-1997
PRIOR APPLICATION DATA: 08/643,798
APPLICATION UNMBER: 08/643,798
FILING DATE: 07-MAY-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                    INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: per
HYPOTHETICAL: NO
ANTI-SENSE: NO
IMMEDIATE SOURCE:
                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                             NAME: Meiklejohn, Ph.D., Anita L. REGISTRATION NUMBER: 35.283 REFERENCE/DOCKET NUMBER: 09404/04. TELECOMMUNICATION INFORMATION: 617/542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/01
FILING DATE: 16-DEC-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
COMPUTER: IBM Compat
OPERATING SYSTEM; Wi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                   TELEPHONE: 617/542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: US
ZIP: 02110-2804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: 225 CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: sir
TOPOLOGY: linear
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   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FastSEQ for Windows Version 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IBM Compatible
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100.0%; Score 47; DB 2;
100.0%; Pred. No. 5e+05;
cive 0; Mismatches
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RESULT 50
US-08-996-139-7
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US-08-792-832A-7
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OPERATING SYSTEM: Apple Operating System 7.5.5
SOPTWARE: Microsoft Word for Power Macintosh 6.0.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/996,139
FILING DATE: 22 DECEMBER 1997
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 60/064,671
FILING DATE: 14 OCTOBER 1997
PRIOR APPLICATION NUMBER: USSN 08/813,509
FILING DATE: 07 MARCH 1997
PRIOR APPLICATION NUMBER: USSN 08/813,509
FILING DATE: 07 MARCH 1997
PRIOR APPLICATION NUMBER: USSN 08/813,509
FILING DATE: 32 DECEMBER 1996
ATTORNEY/ACENT INFORMATION:
NUMBER: DSYMBER: USSN 08/772,330
FILING DATE: 23 DECEMBER 1996
ATTORNEY/ACENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 7, Application US/08996139 Patent No. 6017729
                                                       Sequence 7, Application US/08792832A Patent No. 6017734 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                       Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (206)587-0430
TELEFAX: (206)233-0644
INFORMATION FOR SEQ ID NO: 7:
                                                                                                                                                                                                                                                                                                                                                          IMMEDIATE SOURCE:
CLONE: FLAG_ peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 98101
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
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APPLICANT: Galibert, Laurent
APPLICANT: Maraskovsky, Eugene
TITLE OF INVENTION: Receptor Activator of NF-kappaB
                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 5.
CTTY: Seattle
   APPLICANT:
                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Perkins, Patricia Anne
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2851-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
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Summers Dr., Max D.
Braunagel Dr., Sharon C.
Hong Dr., Tao
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Pred. No.
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FITLE OF INVENTION: LIPOCALIN HOMOLOG
FILE REFERENCE: 97-24
CURRENT APPLICATION NUMBER: US/09/130,663A
CURRENT FILING DATE: 1998-08-05
EARLIER APPLICATION NUMBER: 60/054,867
EARLIER FILING DATE: 1997-08-06
NUMBER OF SEQ ID NOS: 30
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 6
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US-09-130-663-6
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                                                    ; OTHER INFORMATION: Flag affinity peptide US-09-130-663-6
                                                                                                                                                                                                                                                                                                                        Sequence 6, Application US/09130663A Patent No. 6020163
GENERAL INFORMATION:
APPLICANT: Conklin, Darrell C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local 9
Query Match
Best Local Similarity
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FILING DATE: 30-JAN-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/000,955
FILING DATE: 07-JUL-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/678,435
APPLICATION NUMBER: US 08/678,435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 512/474-7577
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 03-JUL-1996
ATTORNEY/AGENT INFORMATION:
NAME: Hibler, David W.
REGISTRATION NUMBER: 41,071
REFERENCE/DOCKET NUMBER: TAMK
TELECOMMUNICATION INFORMATION:
                                                                                      ORGANISM: Artificial Sequence FEATURE:
                                                                                                                           TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOTTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy
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                                                                                                                                                ENGTH: 8
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ZIP: 77210
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
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LIPOCALIN HOMOLOG
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 Score 47;
Pred. No.
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   DB 2;
5e+05;
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US-09-081-180-16

STREET: 16.

COUNTRY:

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Sequence 16, Application US/09040786 Patent No. 6025197
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 60/041,263
FILLING DATE: MARCH 19, 1997
ATTORNEY/AGENT INFORMATION:
NAME: Lingenfelter, Susan E
REGISTRATION NUMBER: 41,156
REFERENCE/DOCKET NUMBER: 97-17C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-442-6675
TELEFAX: 206-442-6678
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS LENGTH: 8 amino acids
                      APPLICANT:
TITLE OF INVENTION: SECRETED SALIVARY ZSIG32
TITLE OF INVENTION: POLYPEPTIDES
NUMBER OF SEQUENCES: 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
                                                                                         APPLICANT: Sheppard, Paul O.
                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: peptide
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CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                          Local Similarity
nes 8; Conserv
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TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/09/081,180 FILING DATE: CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ITLE OF INVENTION:
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1201 Eastlake Ave. E.
                                                                                                                                                                                                                                                                                                          Conservative
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100.0%; Pred. No.
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Best Local Similarity
"herhes 8; Conserve
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US-08-817-145-17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Patent No. 6025329
                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: UTSUMI, Jun
APPLICANT: SUDO, Tetsuo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:

PRIOR APPLICATION:

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 60/041,26:

PRIOR APPLICATION NUMBER: 997

ATTORNEY/AGENT INFORMATION:

NAME: Lingenfelter, Susan E

REGISTRATION NUMBER: 41,156

REFERENCE/DOCKET NUMBER: 97-1:

TELECOMMUNICATION INFORMATION:

TELEPHONE: 206-442-6675
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
APPLICANT:
                                                                                                  COUNTRY: USA
ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                               SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/817,149
FILING DATE: 02-JUL-1997
                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: THERAPEUTITLE OF INVENTION: DISEASES NUMBER OF SEQUENCES: 19
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                ADDRESSEE: Birch, Street: P.O. Box 74'
CITY: Falls Church
STATE: VA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/040,786
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STREET: 1201 E
CITY: Seattle
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STRANDEDNESS: single
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TELEFAX: 206-442-6678
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                                                                                                                                                                                                                                                  E: Birch, Stewart, Kolasch & Birch, LLP
P.O. Box 747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          8 amino acids
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                                                                                                                                                                                                                                                                                                                                                        TANAKA, Yasuhiko
MATSUI, Mizuo
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01 Eastlake Ave.
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                                                                                     Version #1.30
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RESULT 54

US-09-040-786-16

Patent No. 6025197 GENERAL INFORMATION:

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Query Match Best Local (Matches

09-081-180-16

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                                                                                                                                                                                                                                                                                                      RESULT 57
US-08-785-247-9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; OTHER INFORMATION: Peptide sequence US-08-307-896-9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: LYCHGAT, Srinivas Ravi
APPLICANT: LYCHGAT, Srinivas Ravi
TITLE OF INVENTION: MUTANT ACTIVATED GSALPHA AND ADENYLYL
TITLE OF INVENTION: CYCLASE 2 FOR USE AS THERAPEUTIC AGENTS
FILE REFERENCE: 29770
CURRENT APPLICATION NUMBER: US/08/307,896C
CURRENT APPLICATION NUMBER: US/08/307,896C
CURRENT FILING DATE: 1994-09-16
NUMBER OF SEQ ID NOS: 9
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 9
LENGTH: 8
                                                                                                                                                                                        Sequence 9, Application US/08785247
PATENT NO. 6640149
GENERAL INFORMATION:
APPLICANT: Kolesnick, Richard N.
APPLICANT: Liu, Jun
APPLICANT: Zhang, Yuhua
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 9, Application US/08307896C Patent No. 6034071
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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Best Local Similarity
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TELEPAX: 703-205-8050
INFORMATION FOR SEQ ID NO: 17:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 28,977
REFERENCE/DOCKET NUMBER: 76
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: li
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                       TITLE OF INVENTION: ASSAY FOR IDENTIFYING AGENTS WHICH ACT ON THE TITLE OF INVENTION: CERAMIDE-ACTIVATED PROTEIN KINASE, KINASE TITLE OF INVENTION: SUPPRESSOR OF RAS, AND METHODS OF USING SAID AGENTS
                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                          NUMBER OF SEQUENCES:
                STATE: N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
                                                   CITY: New York
                                                                    STREET:
                                                                                      ADDRESSEE:
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10036
                                                                                                                                                                                                                                                                    , Application US/08785247 6040149
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                New York
Y: U.S.A.
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                                                                  E: Cooper & Dunham LLP
1185 Avenue of the Americas
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 47; DB 2; Length 8; 100.0%; Pred. No. 5e+05;
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Pred. No. 5e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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US-08-785-247-9
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Patent No. 6040437
GENERAL INFORMATION:
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Best Local Similarity
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TELEPHONE: 212-278-0400
TELEPAX: 212-881-0526
INFORMATION FOR SEQ ID NO: 9:
NAME: Dreger, Ginger R.
REGISTRATION NUMBER: 33,055
REFERENCE/DOCKET NUMBER: P106
TELECOMPUNICATION INFORMATION:
TELEPHONE: 650/225-3216
TELEFAX: 650/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/938,830
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HYPOTHETICAL:
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NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
COMPUTER: IBM PC com
                                                                                             CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/798419
FILING DATE: 07-FEB-1997
ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Ginger R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Lasky, Laurence A.
APPLICANT: Dowbenko, Donald J.
TITLE OF INVENTION: Tyrosine Phosphorylated Cleavage
TITLE OF INVENTION: Furrow-Associated Proteins (PSTP:
NUMBER OF SEQUENCES: 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 1 DNA Way
CITY: South San Fi
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/785,247 FILING DATE:
                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Genentech, Inc.
STREET: 1 DNA Way
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ENGTH:
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RESULT 60
US-08-819-177-14
; Sequence 14, Application US/08819177
; Patent No. 6043083
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US-08-828-741B-9
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                                                                                                                                                           Query Match
Best Local S
Matches 8
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TELEPHONE: (516) 742-4343
TELEFAX: (516) 742-4366
TELEX: 230 901 SANS UR
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
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APPLICANT: Koentg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION DATA:
                                                                                                                                                                                                                                     TOPOLOGY: 11
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: DiGiglio, Frank S.
REGISTRATION NUMBER: 31,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Tarlinton, David M.
APPLICANT: Treutlein, Herbert R.
TITLE OF INVENTION: CATALYTIC ANTIBODIES AND A METHOD OF
TITLE OF INVENTION: PRODUCING SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                               LENGTH: 8 amino acids
TYPE: amino acid
STRANGENESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/828,741B
FILING DATE: 26-MAR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Garden City
STATE: New York
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8; Conserv
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Amino Acid
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                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                  DNA (genomic)
                                                                                                                                                                      100.0%; Score 47; DB 2; Length 8; 100.0%; Pred. No. 5e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                          31,346
BER: 10591
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Pred. No. 5e+05;
                                                                                                                                                         Mismatches
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US-09-143-470-14
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; OTHER INFORMATION: Flag eptiope
US-08-819-177-14
                                                                                                                                        APPLICANT: Pan, Yang
TITLE OF INVENTION: NEUROTACTIN AND USES THEREFOR
FILE REFERENCE: 09404/049001
CURRENT APPLICATION NUMBER: US/09/143,470
CURRENT FILING DATE: 1998-08-28
NUMBER OF SEQ ID NOS: 20
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 14
                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
 Matches
                  Query Match
Best Local :
                                                                                                                                                                                                                                                                                                        Patent No. 6043086
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 617-542-8906
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                       ORGANISM: Homo sapiens
                                                                                                          TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                         LENGTH: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/819,177
FILING DATE: 28 April 1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compati
OPERATING SYSTEM: Win
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Davis, Roger J.

APPLICANT: Dickens, Martin
TITLE OF INVENTION: INHIBITORS OF THE JNK SIGNAL
TITLE OF INVENTION: TRANSDUCTION PATHWAY AND METHODS OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: 617-542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADUNE
STREET: 222
STREET: 422
                y Match 100.0%; Score 47; DB 2; Local Similarity 100.0%; Pred. No. 5e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Fasse, J. Peter
REGISTRATION NUMBER: 32
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                amino acid
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Conservative
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YSTEM: Windows95
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 Mismatches
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                                   DB 2;
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                                 Length 8;
Indels
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Sequence 2, Application US/09130225
; Betent No. 6057155
; Betent No. 6057150
; BAPPLICANT: Wickham, Thomas J.
APPLICANT: Roelvink, Petrus W.
APPLICANT: Kovesdi, Ince
TITLE OF INVENTION: TARGETING ADE
TITLE OF INVENTION: CONSTRAINED P
NUMBER OF SEQUENCES: 80
CORRESPONDENCE ADDRESS:
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US-08-950-720A-8
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Patent No. 6046028
                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                  Query Match
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APPLICANT: Conkli
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Sawislak, Deborah A REGISTRATION NUMBER: 37,438 REFERENCE/DOCKET NUMBER: 96-TELECOMMUNICATION INFORMATION: TELEPHONE: 206-442-6672
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: PastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskett
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APPLICANT: Jaspers, Stephen R.
TITLE OF INVENTION: INSULIN HOMOLOG
NUMBER OF SEQUENCES: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: 206-442-6678
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
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   ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8 amino acids
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                                                                                                                                                                                                                                                                                                                                     Conservative
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Leydig, Voit & Mayer, Ltd
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                                                   TARGETING ADENOVIRUS WITH USE OF CONSTRAINED PEPTIDE MOTIFS
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RESULT 64
US-09-022-255-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 3, Application US/09022255
CLASSIFICATION:
PRIOR APPLICATION NUMBER: USSN 08/620,694
APPLICATION NUMBER: USSN 08/620,694
FILLING DATE: 21 MARCH 1996
APPLICATION NUMBER: USSN 08/538,765
FILLING DATE: 7 AUGUST 1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 08/410,535
FILING DATE: 23 MARCH 1995
ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia Anne
REGISTRATION NUMBER: 34,695
REFERENCE/DOCKET NUMBER: 2617-B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Yao, Zhengbin
APPLICANT: Spriggs, Melanie
APPLICANT: Fanslow, William
TITLE OF INVENTION: NO. 6072
NUMBER OF SEQUENCES: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 8 FILING DATE: 21-AUG-1996 INFORMATION FOR SEQ ID NO: 2
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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APPLICATION NUMBER: US/09/130,225
                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Power Macintosh
OPERATING SYSTEM: Apple Operating System 7.5.5
SOFTWARE: Microsoft Word for Apple, Version 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/022,255
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LENGTH: 9 amino acids
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OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                        FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 51 Uni
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TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE
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                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP:
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RESULT 65
US-09-022-696-3
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Best Local Similarity 100.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Yao, Zhengbin
APPLICANT: Spriggs, Melanie
APPLICANT: Fanslow, William
TITLE OF INVENTION: No. 6072037el Receptor That Binds IL-17
NUMBER OF SEQUENCES: 10
TYPE: amino acid
STRANDEDNESS: not rele
TOPOLOGY: linear
MOLECULE TYPE: peptide
IMMEDIATE SOURCE:
CLONE: FLAG peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (206)
INFORMATION FOR SEQ ID NO:
                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                          NAME: PERKINS, PALTICIA Anne
REGISTRATION NUMBER: 34,695
REFERENCE/DOCKET NUMBER: 2617-B
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Power Macintosh
OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM 7.5.5
SOFTWARE: Microsoft Word for Apple, Version 6.0
                                                                                                                                                                                                                                                                      CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION UNMBER: USSN 08
FILING DATE: 23 MARCH 1995
                                                                                                                                                                                                                                         CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA: APPLICATION NUMBER: US
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IMMEDIATE SOURCE:
CLONE: FLAG_ peptide
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STRANDEDNESS: not
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                                                                                                    ENGTH:
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                                                                                                  8 amino acids
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                                                                 not relevant
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US-08-883-036A-5
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US-08-883-036A-5
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                Query Match
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Best Local Similarity
Matches 8; Conserv
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APPLICANT: Rauch, Charles
APPLICANT: Walczak, Henning
TITLE OF INVENTION: Receptor That Binds TRAIL
NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kathryn A. Anderson, Immunex Corporation
                                                                                                                                                                                                                                            TELEX: 756822
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/
FILING DATE: 26-UN-1997
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US --L
FILING DATE: 04-UN-1997
PRIOR APPLICATION DATA:
APPLICATION UMBER: US 08/
FILING DATE: 28-MAR-1997
PRIOR APPLICATION DATA:
                                                                TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
ANTI-SENSE: NO
IMMEDIATE SOURCE:
CLONE: FLAG peptide
                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 2625-D TELECOMMUNICATION INFORMATION: TELEPHONE: (206) 587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 1
FILING DATE: 12-MAR-:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: Apple Power Macintosh
OPERATING SYSTEM: Macintosh 7.6
SOFTWARE: Microsoft word, Version 6.0.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 13-FEB-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                       TELEFAX: 12.
TELEFAX: 756822
                                                                                                                                                                         LENGTH: 8 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                       REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                           NAME:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 51 University Street CITY: Seattle,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                        Anderson, Kathryn A.
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                                                                                                                                                                                                                                                                                 (206) 233-0644
                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-MAR-1997
                                                                                                                                      peptide
100.0%; Score 47; DB 2; 100.0%; Pred. No. 5e+05;
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                Length 8;
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RESULT 67

US-09-030-613-35

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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Blorkiewicz, Robert Z.
APPLICANT: Baird, J. Andrew
TITLE OF INVENTION: INHIBITORS OF LEADERLESS PROTEIN EXPORT
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 35, Application US/09030613 Patent No. 6083706
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/030,613
FILING DATE: 25-FEB-1998
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: NO. 6083706tenburg Ph.D., Carol
REGISTRATION NUMBER: 39,317
REFERENCE/DOCKET NUMBER: 39,317
REFERENCE/DOCKET NUMBER: 760100.418C1
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Sheppard, Paul O.
APPLICANT: Grossmann, Angelika
TITLE OF INVENTION: NOVEL TUMOR ANTIGENS
NUMBER OF SEQUENCES: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 35:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                        STREET: Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 6300 CCITY: Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
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                                                                                               COUNTRY:
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                                                                                                                                                                                 ADDRESSEE:
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                                                                                                                                                            E: ZymoGenetics, Inc.
1201 Eastlake Avenue East
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
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US-08-566-190-20
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; MOLECULE TYPE: US-08-566-190-20
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Best Local Similarity
Matches 8; Conserv
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SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                 TELEFAX: (404) 873-8795
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                   CLASSIFICATION: 514
ATTORINEY/AGENT INFORMATION:
NAME: Paber, Patrea L.
REGISTRATION NUMBER: 31,284
REFERENCE/DOCKET NUMBER: YU11
TELECOMMUNICATION INFORMATION:
TELEPHONE: (404) 873-8794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Sawislak, Deborah A REGISTRATION NUMBER: 37,438 REFERENCE/DOCKET NUMBER: 97-TELECOMMUNICATION INFORMATION: TELEPHONE: 206-442-6672
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,569
                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
ZIP: 30309-3450
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Warren, TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 200
STREET: Atlanta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: si
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                                TYPE: amino acid
STRANDEDNESS: sir
TOPOLOGY: linear
                                                                                                                                                                                                                                                                         APPLICATION NUMBER: FILING DATE: 1-DEC-
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                                                                                   LENGTH:
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                                                                                   8 amino acids

    Patrea L. Pabst
    2800 One Atlantic Center, 1201 West Peachtree Street

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ilarity 100.0%;
Conservative (
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RNA Polymerase II Peptides and Methods of Use
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Pred. No. 5e+05;
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RESULT 71
US-09-022-253-3
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US-09-110-618-5
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      Sequence 3, Application US/09022253
Patent No. 6096305
GENERAL INFORMATION:
APPLICANT: Yao, Zhengbin
APPLICANT: Spriggs, Melanie
                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 5, Application US/09110618 Patent No. 6090918
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Anderson, Kathryn A.
REGISTRATION NUMBER: 32,172
REFERENCE/DOCKET NUMBER: 2619
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 587-0430
TELEPAX: (206) 233-0644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (206) 233-06
TELEX: 756822
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                             IMMEDIATE SOURCE:
CLONE: FLAG peptide
                                                                                                                                                                                                                                                                                                                                 HYPOTHETICAL: NO ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/604,333
FILING DATE: 21-FEB-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple 7.1
SOFTWARE: Microsoft Word, Version 5.1a
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kathryn A. Anderson, Immunex Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Parnet, Patricia et al. TITLE OF INVENTION: Receptor Desig
                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 8 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 51 Uni
                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/09/110,618
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                                                                                                                                                                              1 DYKDDDDK 8
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Yao, Zhengpın
Spriggs, Melanie
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                                                                                                                                                                                                                                                                                                                                                                                 linear
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                                                                                                                                                                                                                                 Score 47;
Pred. No. 5
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                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                Length 8;
                                                                                                                                                                                                                     Indels
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                                                                                                                                                                                                                                              RESULT 72
US-09-022-260-3
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Best Local Similarity
Matches 8; Conserva
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                                                                                                                                                                                                                   Sequence 3, Application US/09022260 Patent No. 6100235
                                                                                                                                                                                                                  Patent No.
                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (206)
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                  APPLICANT: Yao, Zhengbin
APPLICANT: Spriggs, Melanie
APPLICANT: Fanslow, Milliam
TITLE OF INVENTION: No. 6100235el Receptor That Binds IL-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                          IMMEDIATE SOURCE:
CLONE: FLAG_ peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 34,695
REFERENCE/DOCKET NUMBER: 26
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/620,694
FILING DATE: 21-MARCH-1996
APPLICATION NUMBER: USSN 08/538,765
FILING DATE: 7 AUGUST 1995
CLASSIFICATION:
                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 23 MARCH
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: Apple Power Macintosh
OPERATING SYSTEM: Apple Operating System 7.5.5
SOFTWARE: Microsoft Word for Apple, Version 6.0.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia Anne
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Fanslow, William
TITLE OF INVENTION: No. 6096305el Receptor That Binds IL-17
NUMBER OF SEQUENCES: 10
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CTTY: Seattle
        ZIP:
                                                STREET: 51 Uni
                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE:
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                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Immunex Corporation STREET: 51 University Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ENGTH:
                                                                                                                                                                                                                                                                                                                                            т рукрарак в
      98101
             WA
USA
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                                                                 E: Immunex Corporation
51 University Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8 amino acids
                                                                                                                                                                                                                                                                                                                                                                              Conservative
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linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          peptide
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100.0%; Pred. No.
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CH 1995
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Length 8; Indels

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Best Local Similarity lov.
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Patent No. 6111073
                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: not rele
TOPOLOGY: linear
MOLECULE TYPE: peptide
IMMEDIATE SOURCE:
CLONE: FLAG peptide
                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPattn (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/020,222
FILING DATE: 06-Feb-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 08/410,535
FILING DATE: 23 MARCH 1995
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Power Macintosh
OPERATING SYSTEM: Apple Operating System 7.5.5
SOFTWARE: Microsoft Word for Apple, Version 6.0.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/022,260
                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Lasky, Laurence A.
APPLICANT: Lasky, Laurence A.
TITLE OF INVENTION: Tyrosine Phosphorylated Cleavage
TITLE OF INVENTION: Furrow-Associated Proteins (PSTP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Perkins, Patricia Anne
REGISTRATION NUMBER: 34,695
REFERENCE/DOCKET NUMBER: 2617-B
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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                                  CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                           STREET: 1 DNA way
CITY: South San Francisco
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ZIP: 94080
APPLICATION NUMBER: 08/
FILING DATE: 02/07/1997
                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
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                                                                                                                                                                                                                                                              California
                                                                                                                                                                                                                                                                                                1 DNA Way
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                   08/798419
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Pred. No. 5e+05;
; Mismatches
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RESULT 74
US-09-053-866-6
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                                                               ; MOLECULE TYPE: US-09-053-866-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 6
Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
Query Match 100.0%; Score 47; DB 2; Best Local Similarity 100.0%; Pred. No. 5e+05; Matches 8; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL
                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Leith, Debra K
REGISTRATION NUMBER: 32,619
REFERENCE/DOCKET NUMBER: 98-1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-442-6674
                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: FASTSEQ for Windows Version CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/053,866 FILING DATE: CLASSTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-3216
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LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 98102
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatik
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Yee, David P.
APPLICANT: Foster, Donald C.
TITLE OF INVENTION: PROTEASE-ACTIVATED RECEPTOR
TITLE OF INVENTION: PAR4 (ZCHEMR2)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Dreger, Ginger
REGISTRATION NUMBER:
                                                                                             LENGTH: 8 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
les 8; Conserv
                                                                                                                                                                                                                                                                                                                           FILING DATE:
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1201 Eastlake Avenue East
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ilarity 100.0%;
Conservative 0
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Presnell, Scott R.
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                                                                                 No. 6111075e
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Pred. No. 5e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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                                   Length 8;
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      Indels
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      Gaps
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GENERAL INFORMATION:

APPLICANT: Crystal, Ronald G.

APPLICANT: Falck-Pedersen, Erik

APPLICANT: Gall, Jason

APPLICANT: Gall, Jason

APPLICANT: Wickham, Thomas J.

TITLE OF INVENTION: CHIMERIC ADENOVIRAL COAT PROTEIN AND

TITLE OF INVENTION: METHODS OF USING SAME

NUMBER OF SEQUENCES: 60

CORRESPONDENCE ADDRESS:

ADDRESSEE: LEYDIG, VOIT & MAYER, LTD.

STREET: TWO PRUDENTIAL PLAZA - 4900

CITY: CHICAGO

CITY: CHICAGO
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                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TOPOLOGY: linear; MOLECULE TYPE: peptide US-08-816-346-50
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                                                                                                                                                                                                                                                   RESULT 76
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US-08-816-346-50
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Best Local S
Matches 8
                                                                                                                                                           Sequence 40, Application US/09071710
Patent No. 6130043
GENERAL INFORMATION:
APPLICANT: BILLING-MEDEL, PATRIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 50, Application US/08816346 Patent No. 6127525
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TELEX: 25-3533
INFORMATION FOR SEQ ID NO:
                                                                                                          APPLICANT:
APPLICANT:
APPLICANT:
                                      APPLICANT:
                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/816,346
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 60601-6780
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 67
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                          Local Similarity 100 
les 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: 312/616-5700
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               f: 8 amino acids amino acid
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COHEN, MAURICE
COLPITTS, TRACEY L.
FRIEDMAN, PAULA N.
GORDON, JULIAN
GRANADOS, EDWARD N.
HODGES, STEVEN C.
KLASS, MICHAEL R.
KRATOCHVIL, JON D.
ROBERTS-RAPP, LISA
                                                                                                                                                               BILLING-MEDEL, PATRICIA
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100.0%; Pri
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US-08-888-429A-16
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                                                                                                                                                                                                                                                                                                                               Sequence 16, Application US/08888429A Patent No. 6136596 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Becker, Cheryl L.
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 608:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
TELEFAX: 847/938-2623
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version
                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                      APPLICANT: Davis, Roger J.
APPLICANT: Whitmarsh, Alan
APPLICANT: Tournier, Cathy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 08/8
FILING DATE: 02-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/071,710
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Abbott Laboratories
STREET: 100 Abbott Park Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                        TITLE OF INVENTION: CYTOKINE-, STRESS-, AND ONCOPROTEIN-TITLE OF INVENTION: ACTIVATED HUMAN PROTEIN KINASE KINASES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Y Match 100.0%;
Local Similarity 100.0%;
                                                                                                ZIP:
                                                                                                                                    STATE: MA
                                                                                                                                                     CITY:
                                                                                                                                                                  STREET:
                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER:
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                                                                                                02110-2804
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                                                                                                                                                 Boston
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                                                                                                                                                                      225 Franklin Street
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STROUPE, STEPHEN D.
VENTION: REAGENTS AND METHODS USEFUL
VENTION: FOR DETECTING DISEASES OF THE PROSTATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IBM Compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              No. 6130043e
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Pred. No. 5e+05;
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CURRENT APPLICATION NUMBER: US/09/432,335
CURRENT FILING DATE: 1999-11-02
EARLIER APPLICATION UNMBER: 09/130,663
EARLIER FILING DATE: 1998-08-06
EARLIER APPLICATION NUMBER: 60/054,867
EARLIER FILING DATE: 1997-08-06
NUMBER OF SEQ ID NOS: 30
SOPTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 6
LENGTH: 8
TYPE: PRT
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US-09-432-335-6
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                                                                               RESULT 79
US-09-079-785-15
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                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: Flag affinity peptide US-09-432-335-6
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Best Local Similarity
Matches 8; Conserv
Sequence 15, Application US/09079785
Patent No. 6143869
GENERAL INFORMATION:
APPLICANT: Goodwin, Raymond G.
APPLICANT: Smith, Craig A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: CONKILL, Darrell C.
TITLE OF INVENTION: LIPOCALIN HOMOLOG
FILE REFERENCE: 97-24
                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 6, Application US/09432335 Patent No. 6143720
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FILING DATE: 19-MAY-1995
ATTORNEY/AGENT INFORMATION:
NAME: Fasse, Peter J.
REGISTRATION NUMBER: 32,983
REFERENCE/DOCKET NUMBER: 07917/0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                           ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
APPLICATION NUMBER: US.
FILING DATE: 07-JUL-19
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: 617/542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 08/5 FILING DATE: 19-SEP-1995 APPLICATION NUMBER: 08/4
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07-JUL-1997
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ÆR: 07917/053001
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Pred. No.
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RESULT 80
US-09-031-168-1
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Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION INVESE: US 907,224
FILING DATE: 01-JUL-1992
PRIOR APPLICATION DATA:
APPLICATION INVESE: US 899,660
FILING DATE: 15-JUN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 892,459
APPLICATION NUMBER: US 892,459
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 02-JUN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 88:
FILING DATE: 26-MAY-1992
ATTORNEY/AGENT INFORMATION:
NAME: Seese, Kathryn A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Armitage, Richard J.
APPLICANT: Gruss, Hans-Jurgen
TITLE OF INVENTION: No. 6143869el Cytokine That Binds
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kathryn A. Seese, Immunex Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple 7.1
SOFTWARE: Microsoft Word, Version 5.1a
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/079,785
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/225,989
FILING DATE: 12 ARRIL 1994
APPLICATION NUMBER: US 07/966,775
FILING DATE: 27-OCT-1992
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 98101
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 32,172
REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
                                                                                                                                                                                                                                          IMMEDIATE SOURCE:
CLONE: FLAG peptide
                                                                                                                                                                                                                                                                              TOPOLOGY: 1
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 1-056822
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                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: (200) 233-0644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: sin
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                                                                                                             1 рукрарак в
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                                                                                                                                                                                                                                                                                                                                                       8 amino acids
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                                                                                                                                                                                                                                                                                            linear
                                                                                                                                                100.0%; Score 47; DB 2; ilarity 100.0%; Pred. No. 5e+05; Conservative 0; Mismatches 0
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02-JUN-1992
                                                                                                                                                                                                                                                                            peptide
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                                                                                                                                                                                      Length 8;
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Sequence 1, Application US/09031168
Patent No. 6150583
GENERAL INFORMATION:
APPLICANT: Stanley B. Prusiner
APPLICANT: Glenn C. Telling

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RESULT 81
US-09-335-411-50
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REFERENCE/DOCKET NUMBER: 0753
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 322-5070
TELEPAX: (415) 854-0875
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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APPLICANT: Michael R.
TITLE OF INVENTION: I
TITLE OF INVENTION: F
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                             tent No.
              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                               APPLICANT: Gall, Jason
APPLICANT: Kovesdi, Imre
APPLICANT: Wickham, Thomas J.
TITLE OF INVENTION: CHIMERIC ADENOVIRAL COAT PROTEIN AND
TITLE OF INVENTION: METHODS OF USING SAME
                                                                                                                                                                                    NUMBER OF SEQUENCES: 6
                                                                                                                                                                                                                                                                                                       APPLICANT: Crystal, Ronald G. APPLICANT: Falck-Pedersen, Erik
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Valeta Gregg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC comparible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
STREET: 2200 Sand Hill Road, Suite 100
CITY: Menlo Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: ASCIII
CURRENT APPLICATION DATA:
                                                                                                    STATE: I
COUNTRY:
                                                                                                                                      STREET:
CITY: C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity 100.0%; Score 47; DB 2; Length 8; les 8; Conservative
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TOPOLOGY: linear
                                                                                                                                                                      ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/09/031,168 FILING DATE:
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                                                                                    60601-6780
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                                                                                                                                                                                                                                                                                                                                                                             Application US/09335411
                                                                                                                                                    E: LEYDIG, VOIT & MAYER, LTD
TWO PRUDENTIAL PLAZA - 4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          8 amino acids
                                                                                                    USA
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Michael R. Sc
PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TRANSGENIC ANIMALS EXPRESSING EPITOPE-TAGGED PROTEINS
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TELEFAX: 31.,
TELEFAX: 25-353
TELEX: 25-353
; INFORMATION FOR SEQ ID NO: 50;
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
TYPE: amino acid
TYPE: 1inear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; TOPOLOGY: linear ; MOLECULE TYPE: peptide US-09-335-411-50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 5, Application US/08883086
Patent No. 6171787
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 8; Conserv
                                                                                                                            NAME: Porembski, Priscilla E
REGISTRATION NUMBER: 33,207
REFERENCE/DOCKET NUMBER: 613.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847-937-0378
TELEFAX: 847-938-2623
                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                            FILING DATE:
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: DOS
SOPTWARE: FastSEQ Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 100...
CITY: Abbott Park
CTATE: IL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: WILEY, STEVEN
TITLE OF INVENTION: MEMB
TITLE OF INVENTION: FOR
NUMBER OF SEQUENCES: 13
                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Abbott Laboratories
STREET: 100 Abbott Park Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
REFERENCE/DOCKET NUMBER: 67
TELECOMMUNICATION INFORMATION:
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PRIOR APPLICATION NUMBER:
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
                                            TYPE: amino acids
                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
                 STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 60064-3500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: 312/0-5700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 DYKDDDDK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 DYKDDDDK 8
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ilarity 100.0%; Pred. No. 5e+05;
Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WILEY, STEVEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     312/616-5600 ,
No. 6171787e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEMBER OF THE TNF FAMILY USEFUL FOR TREATMENT AND DIAGNOSIS OF DISEASE
                                                                                                                                                                                                                                                                                                                            US/08/883,086
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US-08-912-276-24
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                                                                                                                    ; TOPOLOGY: US-08-912-276-24
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GENERAL INFORMATION:
APPLICANT: Billin
APPLICANT: Cohen,
                                                       Query Match
Best Local Similarity
Matches 8; Conserv
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                                                                                                                                              INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                              FILING DATE:

ATTORNEY/AGENT INFORMATION:

NAME: Becker, Cheryl L

REGISTRATION UNMBER: 35,441

REFERENCE/DOCKET NUMBER: 5972

TELECOMMUNICATION INFORMATION:

TELEPHONE: 847/935-1729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/912,276
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: I
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                                                                                                                                                                                                                                      TELEFAX: 847/938-2623
                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
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                    DYKDDDDK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3: Abbott Laboratories
100 Abbott Park Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Roberts-Rapp, Lisa
Russell, John C.
Stroupe, Steven D.
VENTION: REAGENTS AND METHODS USEFUL
VENTION: FOR DETECTING DISEASES OF THE BREAST
VENCENCES: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        USA
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linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cohen, Maurice
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Billing-Medel, Patricia A.
                                                      100.0%; Score 47; DB 2; llarity 100.0%; Pred. No. 5e+05; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gordon, Julian
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Pred. No. 5e+05;
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                                                                                     DB 2; Length 8;
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RESULT 84
US-09-046-158A-20
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US-09-022-259-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIAN PC-DOS/MS-DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/046,158A
FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: DATNLEY Jr., James D.
REGISTRATION NUMBER: 33,673
TELECOMMUNICATION INFORMATION:
TELEPHONE: 616/833-2210
                                                                                                                                                                                                                                                Sequence 3, Application US/09022259
Patent No. 6191104
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 8; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Roberds, Steven L.
APPLICANT: Kaytes, Paul S.
TITLE OF INVENTION: METHOD FOR IDENTIFYING INHIBITORS
TITLE OF INVENTION: JAK2/CYTOKINE RECEPTOR BINDING
NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
ZIP: 98101
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                          APPLICANT: Yao, Zhengbin
APPLICANT: Spriggs, Melanie
APPLICANT: Fanslow, William
TITLE OF INVENTION: No. 6191104el Receptor That
NUMBER OF SEQUENCES: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS
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                                                                      STREET: 51 Unition CITY: Seattle STATE: WA
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COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acids
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                                   COUNTRY: USA
ZIP: 98101
                                                                                                                          ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                   1 DYKDDDDK 8
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                                                                                                          51 University Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      616/833-8897
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linear
                                                                                                                            Immunex Corporation
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Pred. No. 5e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 2;
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                                                                                                                                                                                    Binds IL-17
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                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 3, Application US/09022257
Patent No. 6197525
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Power Macintosh
OPERATING SYSTEM: Apple Operating System 7.5.5
SOFTWARE: Microsoft Word for Apple, Version 6.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                              APPLICANT: Yao, Zhengbin
APPLICANT: Spriggs, Melanie
APPLICANT: Fanslow, William
TITLE OF INVENTION: No. 6197525el Receptor That Binds IL-17
NUMBER OF SEQUENCES: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia Anne
REGISTRATION UMMBER: 34,695
REFERENCE/DOCKET NUMBER: 261
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 08
FILING DATE: 23 MARCH 1995
CLASSIFICATION:
                                                              PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER: 08/620,694
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OPERATING SYSTEM: Apple Operating System 7.5.5
SOFTWARE: Microsoft Word for Apple, Version 6.0.1
CURRENT APPLICATION DATA:
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               FILING DATE:
CLASSIFICATION:
                                             APPLICATION NUMBER:
                                                                           FILING DATE:
CLASSIFICATION:
                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
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APPLICATION DATA:
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51 University Street
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                                           08/620,694
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Pred. No. 5e+05;
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RESULT 88
US-09-105-343A-5
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                                                                                                                                                                                      Query Match
Best Local S
Matches 8
Sequence 5, Application US/09105343A Patent No. 6207642 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 17
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 16, Application US/09153804 Patent No. 6207380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Jon D. Kratochvil
APPLICANT: Eric Russell
APPLICANT: Stephen D. Stroupe
TITLE OF INVENTION: Reagents and Methods Useful for Detecting Diseases of the Urinar
TITLE OF INVENTION: Tract
FILE REFERENCE: 6180.US.01
CURRENT APPLICATION NUMBER: US/09/153,804
CURRENT FILING DATE: 1998-09-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANCE
TOPOLOGY: linear
MOLECULE TYPE: peptide
IMMEDIATE SOURCE:
THONE: FLAG_ peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Julian Gordon
APPLICANT: Steven C. Hodges
APPLICANT: Michael R. Klass
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Maurice Cohen
APPLICANT: Paula N. Friedman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Patricia Billing-Medel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                      OTHER INFORMATION: Affinity purification system recognition site
                                                                                                                                                                                                                                                                                             TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                ENGTH: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 26
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
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                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Perkins, Patricia Anne REGISTRATION NUMBER: 34,695
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: FILING DATE: 23 MAR CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ENGTH:
                                                                                                                                                       1 DYKODDDK 8
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8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Score 47; DB 2; ilarity 100.0%; Pred. No. 5e+05; Conservative 0; Mismatches
                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 not relevant
                                                                                                                                                                                      100.0%; Score 47; DB 2;
100.0%; Pred. No. 5e+05;
tive 0; Mismatches
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PPLICANT: WILEY, S.R.

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; ORGANISM: Mus musculus 
US-08-693-211-1
                                                                                                                                                                                                                          Sequence 1, Application US/08693211
Patent No. 6218516
GENERAL INFORMATION:
APPLICANT: Wilks, Andrew
APPLICANT: Stacker, Steven
APPLICANT: Oelrichs, Robert
TITLE OF INVENTION: Antibodies Specific For the Extracellular Domain of NYK/FLK-1
TITLE OF INVENTION: And uses Thereof
FILE REFERENCE: LUD 5347
                                                                          NUMBER OF SEQ ID NOS: SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                              CURRENT APPLICATION NUMBER: US/08/693,211
CURRENT FILING DATE: 1996-09-30
PRIOR APPLICATION NUMBER: PCT/US95/01727
PRIOR FILING DATE: 1995-02-09
PRIOR APPLICATION NUMBER: PM3793
PRIOR APPLICATION NUMBER: PM3793
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: BECKER, Cheryl L.
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 6448
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847-935-1729
TELEPAX: 847-938-2623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                PRIOR FILING DATE: 1994-02-10
                                      TYPE: PRT
                                                          LENGTH: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US98/02859
FILING DATE: 12-FEB-1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
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ADDRESSEE: Abbott Laboratories
STREET: 100 Abbott Park Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: MEMBER OF THE TITLE OF INVENTION: FOR TREATMENT NUMBER OF SEQUENCES: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: No. 6207642e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity hes 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/0 FILING DATE: 12-FEB-1998 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Abbott Park
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Pred. No.
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RESULT 90
US-08-944-483-75
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US-08-944-483-75
                                                             Best Loc
Matches
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Best Local Similarity
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Patent No. 6232456
                                                                                           Query Match
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                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatibl
OPERATING SYSTEM: DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                   NAME: Becker, Cheryl L.
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 61
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/335-1729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: RUSSELL, JOHN C.
APPLICANT: STEWART, KENT D.
APPLICANT: STEWART, KENT D.
APPLICANT: STROUPE, STEVEN D.
TITLE OF INVENTION: NOVEL SERINE PROTEASE REAGENTS
TITLE OF INVENTION: AND METHODS USEFUL FOR DETECTING
TITLE OF INVENTION: OF THE PROSTATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
APPLICANT:
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ATTORNEY/AGENT INFORMATION:
NAME: Becker, Cheryl L.
                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: FastSEQ for Windows Version CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/944,483
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
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                                                                                                                                                          LENGTH: 8 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Abbott Laboratories STREET: 100 Abbott Park Road
                                                                            Local
                                                                                                                                                                                                                                                         TELEFAX: 847/938-2623
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                                                             Similarity
8; Conserv
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nilarity 100.0%;
Conservative 0;
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KLASS, MICHAEL R.
                                                                                                                                             linear
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                                                               Score 47; DB
Pred. No. 5e+
0; Mismatches
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Pred. No. 5e+05;
Mismatches
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                                                                                             Length 8;
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RESULT 91 US-09-227-717-5 ; Sequence 5, Application US/09227717

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                                                                                                                                                             PATENT NO. 6242213

GENERAL INFORMATION:
APPLICANT: Anderson, Dirk M.
APPLICANT: Galibert, Laurent
APPLICANT: Maraskovsky, Eugene
TITLE OF INVENTION: Ligand for Receptor Activator of NF-kappaB
NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 8; Conserv
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO:
 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Power Macintosh
OPERATING SYSTEM: Apple Operati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: MCMasters, David D.
REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 690
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
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ADDRESSEE: Seed and Berry
STREET: 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: DeSouza, Errol B.
TITLE OF INVENTION: INTERLEUKIN-1 TYPE 3 RECEPTORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Lovenberg, Timothy W. APPLICANT: Oltersdorf, Tilman
                                                                         COUNTRY: US
ZIP: 98101
                                                                                                                            STREET: 51 Uni
CITY: Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
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                                                                                          USA
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Clevenger, William R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
Apple Operating System 7.5.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 47; DB 2; 100.0%; Pred. No. 5e+05;
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Sequence 40, App. ...
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (206)233-0644
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                               APPLICANT:
ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                           NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: peptide IMMEDIATE SOURCE: CLONE: FLAG_ peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia Anne
REGISTRATION NUMBER: 34,693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 08/772,330
FILING DATE: 23 DECEMBER 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 08/813,509
FILING DATE: 07 MARCH 1997
                                                                                                                                                                 TITLE OF INVENTION:
                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                 APPLICANT:
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                                                                                                                                                                               FITLE OF INVENTION:
                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                 COUNTRY:
                                                                                                                 ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS: not
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2852-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                              Application US/09525397
                                                                                                100 Abbott Park Road
                                                                                                                                                                                                                                                                                          COHEN, MAURICE
COLPITTS, TRACEY L.
FRIEDMAN, PAULA N.
GORDON, JULIAN
GRANADOS, EDWARD N.
                                                USA
                                                                                                                                                                             ROBERTS-RAPP, LISA
RUSSELL, JOHN C.
STROUPE, STEPHEN D.
NVENTION: REAGENTS AN
                                                                                                                                                                                                                            HODGES, STEVEN C. KLASS, MICHAEL R. KRATOCHVIL, JON D. ROBERTS-RAPP, LISA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (206) 233-0644
                                                                                                                                                                                                                                                                                                                                                                              BILLING-MEDEL, PATRICIA
                                                                                                            Abbott Laboratories
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Microsoft Word for Power Macintosh 6.0.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TUMBER: US/08/995,659
22 DECEMBER 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14 OCTOBER 1997
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                                                                                                                                                                 FOR DETECTING
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CTING DISEASES OF THE PROSTATE
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RESULT 94
US-09-320-774-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Jarvik
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Becker, Cheryl L.
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 6083
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
TELEFAX: 847/938-2623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
                                      REGISTRATION NUMBER: 35,461
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 654-2428
INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                        CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of COMPUTER: IBM PC computer:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Jarvik, Jonathan
TITLE OF INVENTION: READING
TITLE OF INVENTION: TAGGING
NUMBER OF SEQUENCES: 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: No. 6252047e
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SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
                                                                                                                ATTORNEY/AGENT INFORMATION: NAME: Brotman, Harris F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                  SEQUENCE CHARACTERISTICS:
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                                                                                          NAME: Brotman, Harris
REGISTRATION NUMBER: 3
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                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: La Jolla
STATE: California
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LENGTH:
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202_Coast Blvd., Suite 111
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  8 amino acids
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RESULT 95
US-09-215-649A-7
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                                                                                                            IMMEDIATE SOURCE:
CLONE: FLAG peptide
SEQUENCE DESCRIPTION: SEQ
US-09-215-649A-7
                                        Best Loc
Matches
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GENERAL INFORMATION:
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Matches 8; Conserv
                                                                         Query Match
                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 7
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                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/996,139

PILING DATE: <Unknown-
APPLICATION WIMBER: USSN 08/813,509

FILING DATE: 07 MARCH 1997

APPLICATION NUMBER: USSN 08/772,330

PILING DATE: 23 DECEMBER 1996

ATTORNEY/AGENT INFORMATION:

NAME: Perkins, Patricia Anne
REGISTRATION NUMBER: 34,693
                                        Local Similarity
les 8; Conserv
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OPERATING SYSTEM: Apple Operating System 7..
SOFTWARE: Microsoft Word for Power Macintosi
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/09/215,649A
FILING DATE: 17-Dec-1998
CLASSIFICATION: Unknown>
                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
TELEFAX: (206)233-0644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 98101
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Maraskovsky, Eugene
TITLE OF INVENTION: Receptor Activator of NF-kappaB
NUMBER OF SEQUENCES: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Anderson, Dirk M.
                                                                                                                                                                                    TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
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DYKDDDDK 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Immunex Corporation, Law Department
STREET: 51 University Street
                                                                                                                                                                                                                   STRANDEDNESS: not relevant
                                                                                                                                                                                                                                      TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
                                                                                                                                                                                                                                                        LENGTH: 8 amino acids
                                    100.0%; Score 47; DB 2; nilarity 100.0%; Pred. No. 5e+05; Conservative 0; Mismatches
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                                                                                                                                                                               peptide
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Pred. No. 5e+05;
Mismatches
                                                                                                                                  ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; OTHER INFORMATION: Description of Artificial Sequence: US-09-053-941-1
                                                                                                                                                                                    SEQ ID NO 7
LENGTH: 8
                                                                                Query Match
Best Local
                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
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Best Local (
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                                                                                                                                    ORGANISM: synthetic
-09-320-424-7
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                                                                                                                                                                                                                                   EARLIER APPLICATION NUMBER: 08/496,632
EARLIER FILING DATE: 1995-06-29
NUMBER OF SEQ ID NOS: 25
                                                                                                                                                                                                                                                                                      EARLIER APPLICATION NUMBER: 08/670,354
EARLIER FILING DATE: 1996-06-25
EARLIER APPLICATION NUMBER: 08/548,368
EARLIER FILING DATE: 1995-11-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Wiley, Steven R.
APPLICANT: Goodwin, Raymond G.
TITLE OF INVENTION: Cytokine that Induces Apoptosis
                                                                                                                                                                                                                        SOFTWARE: PatentIn Ver.
                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/320,424
CURRENT FILING DATE: 1999-05-26
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APPLICANT: KOPROWSKI, HILARY
TITLE OF INVENTION: CHIMERIC VIRAL PROTEINS
FILE REFERENCE: Chimeric Viral Proteins
CURRENT APPLICATION NUMBER: US/09/053,941
CURRENT FILING DATE: 1998-04-02
                                                                                                                                                                                                                                                                                                                                                             EARLIER APPLICATION NUMBER: 09/048,641
                                                                                                                                                                                                                                                                                                                                                                                               EARLIER FILING DATE: 1998-11-10
                                                                                                                                                                                                                                                                                                                                                                                                              EARLIER APPLICATION NUMBER: 09/190,046
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 2835-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EARLIER APPLICATION NUMBER: 60/043,380 EARLIER FILING DATE: 1999-04-03 NUMBER OF SEQ ID NOS: 25
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Pred. No. 5e+
0; Mismatches
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                                                                                          DB 2; Length 8;
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RESULT 98

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SOFTWARE:
SEQ ID NO 6
                                                                                                                               APPLICANT: I
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Patent No. 6300065
GENERAL INFORMATION:
APPLICANT: Kieke, et al.
                                              APPLICANT: Gite, Sadanand
APPLICANT: Olejnik, Jerzy
TITLE OF INVENTION: N-Terminal and C-Te
FILE REFERENCE: AMBER-03879
CURRENT APPLICATION NUMBER: US/09/382,950
CURRENT FILING DATE: 1999-08-25
                                                                                                                                                                                             Sequence 6, Application US/09382950 Patent No. 6303337
                                                                                                                                                                                                                                                                                                                                                                               Matches
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Best Local 9
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Sakano, Seiji
APPLICANT: Icò, Akira
APPLICANT: Icò, Akira
TITLE OF INVENTION: DIFFERENTIATION-SUPPRESSIVE POLYPEPTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 8
TYPE: PRT
ORGANISM: Unknown
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               NUMBER OF SEQ ID NOS:
SOFTWARE: Patentin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Yeast Cell Surface Display of Proteins and Uses Thereof FILE REFERENCE: D6061CIP2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: KP-8576
CURRENT APPLICATION NUMBER: US/09/214,278
CURRENT FILING DATE: 1999-01-26
NUMBER OF SEQ ID NOS: 32
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TYPE: PRT
ORGANISM: Artificial Sequence
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| COMPUTER READABLE FORM: MEDIUM TYPE: Diskett MEDIUM TYPE: Diskett COMPUTER: IBM Compat COMPUTER: IBM Compat COMPUTER: TAM COMPAT COMPUTER: PASTSEM: WISHER COMPUTER: PASTSEM: WISHER COMPUTER: PASTSEM: PASTSEM: WISHER COMPUTER: PASTSEM: APPLICATION NUMBER: FILING DATE: PASTSEMICATION NUMBER: ATTORNEY/AGENT INFORMAT NAME: GOLIET, Minmi ACT REFERENCE/DOCKET NUMBER: REFERENCE/DOCKET NUMBER: | RESULT 1 US-08-971-317A-9 ; Sequence 9, Application ; Publication No. US200100 ; GENERAL INFORMATION: APPLICANT: Wiley, St TITLE OF INVENTION: ; TITLE OF INVENTION: ; NUMBER OF SEQUENCES: ; CORRESPONDENCE ADDRES ; CORRESPONDENCE ADDRES ; CORTEST: 100 Abbott ; STATE: IL ; COUNTRY: USA | 470 470 471 471 471 471 471 471 471 471 | 467 47 100.0 8 468 47 100.0 8 468 47 100.0 8 |
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TOPOLOGY: US-08-926-626-12
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 Query Match
Best Local Similarity
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                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Becker, Cheryl L
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 6158
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
                                                                                                      INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS LENGTH: 8 amino acids
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                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: LIM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION NUMBER: US/08/926,626
APPLICATION NUMBER: US/08/926,626
                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
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                                                                          TYPE: amino acid
STRANDEDNESS: sin
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nes 8; Conserv
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TYPE: amino acid
STRANDEDNESS: single
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(847) 938-2623
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STROUPE, STEVEN D.
PERCENTS AND METHODS USEFUL
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 Score 47;
Pred. No.
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Pred. No. 1.9e+06;
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US-09-042-643-3
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PRIOR FILING DATE: 2000-01-11
PRIOR PELICATION NUMBER: US 08/991,426
PRIOR FILING DATE: 1997-12-16
PRIOR APPLICATION NUMBER: US 08/851,160
PRIOR APPLICATION NUMBER: US 08/643,798
PRIOR APPLICATION NUMBER: US 08/643,798
PRIOR FILING DATE: 1996-05-07
                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/042,643
CURRENT FILING DATE: 1998-03-17
NUMBER OF SEQ ID NOS: 5
SOFTWARE: PatentIn version 3.1
SEQ ID NO 3
                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:

APPLICANT: DAUNERT, Sylvia
APPLICANT: LEWIS, Jennifer C
APPLICANT: HERNANDEZ, Emily
TITLE OF INVENTION: QUANTITATIVE BINDING ASSAYS USING GREEN FLUORESCENT PROTEIN AS
FILE REFERENCE: 3798-0102P
                                     Matches
                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 3, Application US/09042643 Publication No. US20010008766A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 14, Applicati
Publication No. US2001
GENERAL INFORMATION:
APPLICANT: Pan, Yang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity 100 Matches 8; Conservative
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                                                                                                                                       LENGTH: 8
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 18
SOFTWARE: FastSEQ for Windows Version 3.0
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TYPE: PRT
ORGANISM: Homo sapiens
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                                     Conservative
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                                 100.0%; Score 47; DB 3;
100.0%; Pred. No. 1.9e+06;
tive 0; Mismatches 0;
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                                                                  Length 8
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                                     Indels
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RESULT 6
US-09-050-516-48
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APPLICANT: Certretti, Douglas
APPLICANT: Reddy, Pranhitha
APPLICANT: Reddy, Pranhitha
TITLE OF INVENTION: POLYNUCLEOTIDES ENCODING CYTOKINE DESIGNATED LERK-5
FILE REFERENCE: 28232
CURRENT APPLICATION NUMBER: US/09/754,105
CURRENT FILING DATE: 2001-01-03
PRIOR APPLICATION NUMBER: 09/329,531
PRIOR APPLICATION NUMBER: 09/329,531
PRIOR FILING DATE: 1999-06-10
NUMBER OF SEQ ID NOS: 3
SOFTWARE: Patentin version 3.0
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ORGANISM: Artificial
PEATURE:
OTHER INFORMATION: Flag peptide
NAME/KEY: Artificial
LOCATION: (1)...(8)
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Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                            APPLICANT: GORDON, J
APPLICANT: GRANADOS,
APPLICANT: HAYDEN, M
APPLICANT: HODGES, S
APPLICANT: KLASS, MI
APPLICANT: KRATOCHT
APPLICANT: ROBERTS-R
APPLICANT: ROBERTS-R
APPLICANT: RUSSELL,
APPLICANT: STROUPE,
TITLE OF INVENTION:
TITLE OF INVENTION:
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APPLICANT:
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              COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/050,516
FILING DATE:
CLASSIFICATION:
                                                                                                                                                  ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSEE: Abbott Laboratories
STREET: 100 Abbott Park Road
CITY: Abbott Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                        COUNTRY: USA
                                                                                                                                                                                                                                  STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION:
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                                                                                                                                                                                                                                                                                                                                      HODGES, STEVVEN C.

KLASS, MICHAEL R.

KRATOCHVIL, JON D.

ROBERTS-RAPP, LISA
RUSSELL, JOHN C.

STROUPE, STEPHEN D.

VENTION: REAGENTS AND METHODS USEFUL

VENTION: FOR DETECTING DISEASES OF THE GASTROINTESTINAL

ROTENCIES. 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GRANADOS, EDWARD N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GORDON, JULIAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COHEN, MAURICE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BILLING-MEDEL, PATRICIA
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Patricia Billing-Medel
APPLICANT: Tracey L. Colpitts
APPLICANT: Tracey L. Colpitts
APPLICANT: Julian Gordon
APPLICANT: Julian Gordon
APPLICANT: John C. Russell
APPLICANT: Schephen D. Stroupe
TITLE OF INVENTION: Reagents and Methods Useful for
TITLE OF INVENTION: Detecting Disease of the Prostate
FILE REFERENCE: 6397.US.01
CURRENT APPLICATION NUMBER: US/09/276,600
CURRENT FILING DATE: 1999-03-25
NUMBER OF SEQ ID NOS: 11
SOFFWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 10
LENGTH: 8
TYDE: DET
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                                                                                                                                                                        RESULT 8
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                                                                      Sequence 12, Application US/09771956
Patent No. US20010031474A1
GENERAL INFORMATION:
APPLICANT: Bennett, Michele
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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Best Local Similarity
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                  APPLICANT: Brodbeck, Robbin
APPLICANT: Krause, James
TITLE OF INVENTION: Chimeric Neuropeptide Y Receptors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
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REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 60
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 8 amino acids TYPE: amino acid STRANDEDNESS: single
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                                                      Bennett, Michele
Brodbeck, Robbin
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                                                                                                                                                                                                                                                                                                             100.0%; Score 47; DB 3;
100.0%; Pred. No. 1.9e+06;
tive 0; Mismatches 0;
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Best Local S
Matches 8
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CURRENT FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 31
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 12
                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/842,38
FILING DATE: 23-APR-1997
ATTORNEY/AGENT INFORMATION:
NAME: Becker, Cheryl L.
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 6084
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
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APPLICANT:
APPLICANT:
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: REAGENTS AND MELAULO COLLUSTITUS OF INVENTION: FOR DETECTING DISEASES OF THE PROSTATE NUMBER OF SEQUENCES: 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
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ZIP: 60064-3500
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                                                                       TELEFAX:
                                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Abbott Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
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COHEN, MAURICE
COLPITTS, TRACEY L.
FRIEDMAN, PAULA N.
GORDON, JULIAN
GORDON, JULIAN
GRANADOS, EDWARD N.
GRANADOS, STEVEN C.
KLASS, MICHAEL R.
KRATOCHVIL, JON D.
ROBERTS-RAPP, LISA
ROSSELL, JOHN C.
STROUPE, STEPHEN D.
INVENTION: REAGENTS AND METHODS USEFUL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3: Abbott Laboratories
100 Abbott Park Road
                                                                     847/938-2623
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Pred. No. 1.9e+06;
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                                                                                                                FILE REFERENCE: 2879-US
CURRENT APPLICATION NUMBER: US/09/835,147
CURRENT FILING DATE: 2001-04-13
PRIOR APPLICATION NUMBER: US 60/104,585
PRIOR FILING DATE: 1998-10-16
PRIOR APPLICATION NUMBER: US 60/107,466
PRIOR FILING DATE: 1998-11-06
PRIOR FILING DATE: 1998-11-06
PRIOR APPLICATION NUMBER: US 60/149,010
PRIOR APPLICATION NUMBER: US 60/149,010
PRIOR FILING DATE: 1999-08-13
PRIOR APPLICATION NUMBER: PCT/US99/22955
PRIOR FILING DATE: 1999-10-13
NUMBER OF SEQ ID NOS: 31
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US-09-785-934-3
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GENERAL INFORMATION:
APPLICANT: Maliszewski, Charles R:
APPLICANT: Gayle III, Richard B.
APPLICANT: Gayle III, Richard B.
                                                                                SOFTWARE: P
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CURRENT APPLICATION NUMBER: US/09/785,934
CURRENT FILING DATE: 2001-02-15
                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
APPLICANT:
APPLICANT:
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Patent No. US20020002277*
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APPLICANT: Gimpel, Steven D.
TITLE OF INVENTION: Inhibitors of Platelet Activation and Recruitment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 09/579,808
PRIOR FILING DATE: 2000-05-26
NUMBER OF SEQ ID NOS: 3
             TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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                                                          LENGTH: 8
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US20020002132A1
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; OTHER INFORMATION: Description of Artificial Sequence: Synthetic; OTHER INFORMATION: peptide US-09-835-147-10
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US-09-728-911-11
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                                                                                                                                                  Sequence 11, Application US/09728911
PATENT NO. US20020012669A1
GENERAL INFORMATION:
APPLICANT: Presnell, Scott R.
APPLICANT: Xu, Wenfeng
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Best Local
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LENGTH: 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE: 2000-03-15
PRIOR APPLICATION NUMBER: 60/215,644
PRIOR FILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: DK PA 2000 00024
PRIOR FILING DATE: 2000-01-10
PRIOR FILING DATE: 2000-03-02
PRIOR FILING DATE: 2000-03-02
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APPLICANT: MIKKELSEN, JAN MOLLER
TITLE OF INVENTION: G-CSF CONJUGATES
FILE REFERENCE: 31-000700US
                                                                          APPLICANT: Xu, Wenfeng
APPLICANT: Kindsvogel, Wayne
APPLICANT: Chen, Zhi
TITLE OF INVENTION: Human Cyt
FILE REFERENCE: 99-93
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PRIOR FILING DATE: 2000-01-14
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FILE REFERENCE: 99-93
CURRENT APPLICATION NUMBER: US/09/728,911
CURRENT FILING DATE: 2000-12-01
PRIOR APPLICATION NUMBER: US 60/169,049
PRIOR FILING DATE: 1999-12-03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence FEATURE:
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8; Conserv
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                                                                                                Human Cytokine Receptor
                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
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Pred. No.
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FILE REFERENCE: D0016.np
CURRENT APPLICATION NUMBER: US/09/813,329
CURRENT FILING DATE: 2001-03-20
PRIOR APPLICATION NUMBER: 60/190,816
PRIOR FILING DATE: 2000-03-21
NUMBER OF SEQ ID NOS: 65
SOFTWARE: PatentIn version 3.0
SEQ ID NO 65
LENGTH: 8
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US-09-813-329-65
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Best Local Similarity
""" hes 8; Conserve
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; ORGANISM: bacteriophage T7
US-09-813-329-65
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                                                                                                                                                                                                                                                                                                                   RESULT 15
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APPLICANT: Bristol-Myers Suibb Company
APPLICANT: INVENTION: No. US20020012968Alel Drosophila Tumor Necrosis Factor Class Mole
TITLE OF INVENTION: Variants Thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: SEQ ID NO 11
                                                                                                                                                                                                                                            Sequence 26, Application US/09099823
Patent No. US20020018990A1
GENERAL INFORMATION:
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PRIOR FILING DATE: 2000-09-13
PRIOR APPLICATION NUMBER: US
PRIOR FILING DATE: 2000-10-31
NUMBER OF SEQ ID NOS: 36
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TYPE: PRT
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APPLICANT:
APPLICANT:
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APPLICANT:
APPLICANT:
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APPLICANT: YU, HONG
TITLE OF INVENTION:
TITLE OF INVENTION:
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               T: YU, HONG INVENTION:
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                                              HODGES, STEVEN C.
KLASS, MICHAEL R.
KRATOCHVIL, JON D.
RUSSELL, JOHN C.
SCHEFFEL, CHRISTI
STROUPE, STEPHEN D.
                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 47; DB 3; ilarity 100.0%; Pred. No. 1.9e+06; Conservative 0; Mismatches 0.
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                                                                                                                                                                               COHEN, MAURICE COLPITTS, TRACEY L. FRIEDMAN, PAULA N.
                                                                                                                                                GORDON, JULIAN
GRANADOS, EDWARD N.
                                                                                                                                                                                                                                  BILLING-MEDEL, PATRICIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 47; DB 3; 100.0%; Pred. No. 1.9e+06;
REAGENTS AND METHODS USEFUL FOR DETECTING DISEASES OF THE
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US-09-790-264-68
; Sequence 68, Application Us; Patent No. US20020028508A1; GENERAL INFORMATION:
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                                    CURRENT APPLICATION NUMBER: US/09/790,264
CURRENT FILING DATE: 2001-02-21
PRIOR APPLICATION NUMBER: US 09/065,661
PRIOR FILING DATE: 1998-04-23
PRIOR FILING DATE: 1999-04-23
PRIOR PPLICATION NUMBER: US 09/298,531
PRIOR APPLICATION NUMBER: US 09/065,363
PRIOR APPLICATION NUMBER: US 09/065,363
PRIOR APPLICATION NUMBER: US 09/065,363
PRIOR FILING DATE: 1998-04-23
PRIOR PILING DATE: 1998-04-23
PRIOR APPLICATION NUMBER: US 09/337,930
PRIOR PILING DATE: 1999-06-22
PRIOR PILING DATE: 1998-06-22
PRIOR FILING DATE: 1998-06-22
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                                                                                                                                                                                                                                                                                                                         APPLICANT: Holtzman, Douglas A.
APPLICANT: Goodearl, Andrew D.J.
APPLICANT: MCCarthy, Sean A.
TITLE OF INVENTION: NOVEL GENES ENCODING PROTEINS HAVING
TITLE OF INVENTION: PROGNASTIC, DIAGNOSTIC, PREVENTIVE,
TITLE OF INVENTION: USES
                                                                                                                                                                                                                                                                                                           FILE REFERENCE: 07334-322001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Becker, Cheryl L.
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 61
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: No. US20020018990Ale
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LENGTH: 8 amino acids
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FILING DATE: 20-JUN-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
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SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Abbott Laboratories
STREET: 100 Abbott Park Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
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STRANDEDNESS: sir
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APPLICATION NUMBER: US 09/363,630 FILING DATE: 1999-07-29
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Pred. No. 1
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RESULT 18
US-09-234-717-24
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                                                   APPLICANT: Julian Gordon
APPLICANT: Edward N. Granados
APPLICANT: Steven C. Hodges
APPLICANT: Michael R. Klass
APPLICANT: John D. Kratochvil
APPLICANT: John C. Russell
APPLICANT: Stephen D. Stroupe
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Best Local S
Matches 8
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APPLICANT: Lohning, Corinna
APPLICANT: Lohning, Corinna
TITLE OF INVENTION: No. US20020034733A1el methods for displaying (poly) peptides/prot
TITLE OF INVENTION: particles via disulfide bonds
FILE REFERENCE: MORPHO/11
CURRENT ENLIGHENCE: MORPHO/11
CURRENT FILING DATE: 2001-03-15
PRIOR APPLICATION NUMBER: EP 99114072.4
PRIOR PELICATION NUMBER: EP 99114072.4
PRIOR PILING DATE: 1999-07-20
                                                                                                                                                                                                                                                                                               Sequence 24, Application US/09234717A Patent No. US20020034740A1
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Best Local :
                                                                                                                                                                                           APPLICANT: Patricia Billing-Medel
APPLICANT: Maurice Cohen
APPLICANT: Tracey L. Colpitts
APPLICANT: Paula N. Friedman
                TITLE OF INVENTION: Reagents and Methods Useful for Detecting Diseases of TITLE OF INVENTION: the Breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: EP 00103551.8 PRIOR FILING DATE: 2000-02-18 NUMBER OF SEQ ID NOS: 41
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NUMBER OF SEQ ID NOS: 68
SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
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TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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REFERENCE: 6450.US.01
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Similarity 100.0%;
8; Conservative 0
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Pred. No. 1.9e+06;
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Pred. No. 1.9e+06;
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                                                                                  RESULT 20
US-09-193-538-22
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                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: Affinity purification system recognition US-09-850-178-18
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CURRENT FILING DATE: 1999-01-21
NUMBER OF SEQ ID NOS: 25
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 24
LENGTH: 8
GENERAL INFORMATION:
APPLICANT: Patricia Billing-Medel
APPLICANT: Maurice Cohen
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                                                    Sequence 22, Application US/09193538A Patent No. US20020037503A1
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Best Local &
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SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/850,178
CURRENT FILING DATE: 2001-05-07
PRIOR APPLICATION NUMBER: US 08/972,376
PRIOR FILING DATE: 1997-11-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Kratochvil, Jon D.
APPLICANT: Roberts-Rapp, Lisa
APPLICANT: Strouge, Stephen D.
APPLICANT: Gordon, Juliano
TITLE OF INVENTION: REAGENTS AND METHODS USEFUL
TITLE OF INVENTION: DETECTING DISEASES OF THE E
FILE REFERENCE: 6251.US.Pl
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Granados, Edward N.
Hodges, Steven C.
Klass, Michael R.
                                                                                                                                                  DYKODDDK 8
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100.0%; Pred. No.
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RESULT 21
US-09-250-883-22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Lisa Roberts-Rapp
APPLICANT: John C. Russell
APPLICANT: Stephen D. Stroupe
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INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
                                                                                                                                                                                                                                     SOFTWARE: FASTSEQ for Windows Version CURRENT APPLICATION DATA:
APPLICATION NUMBERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ENGTH:
                                                NAME: Becker, Cheryl L
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 61
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Abbott Laboratories
STREET: 100 Abbott Park Road
                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Russell, John APPLICANT: Colpitts, Tra
                                                                                                                                  ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                STREET: 100 ._
CTTY: Abbott Park
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                                                                                                                                                                                                       FILING DATE:
CLASSIFICATION:
                                TELEFAX:
                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
ZIP: 60064-3500
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: Edward N. Granados
: Steven C. Hodges
: Michael R. Klass
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ilarity 100.0%; Pred. No. 1.:
Conservative 0; Mismatches
                                  847/938-2623
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23
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                                                                                      6131.US
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ches 0;
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US-09-096-259-32
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: Flag peptide sequence US-09-735-368-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-250-883-22
                                                                                                                                                                                                                                                                               Sequence 32,
Patent No. U
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Best Local :
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Best Local Similarity
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                                                                                                                                                                                                                                                                     GENERAL INFORMATION
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CURRENT FILING DATE: 2000-12-12
PRIOR APPLICATION NUMBER: US 60/171,258
PRIOR FILING DATE: 1999-12-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                       APPLICANT: HODGES, S
APPLICANT: KLASS, MI
APPLICANT: KRATTOCHVI
APPLICANT: ROBERTS-R
APPLICANT: RUSSELL,
APPLICANT: STROUPE,
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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ITLE OF INVENTION: RING FINGER PROTEIN ZAPOP2
                             CORRESPONDENCE ADDRESS:
                                         NUMBER OF SEQUENCES:
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 STREET:
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STRANDEDNESS: sir
TOPOLOGY: linear
                DDRESSEE:
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8; Conserv
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                                                                                 HODGES, STEVEN C.
KLASS, MICHAEL R.
KRATOCHVIL, JON D.
ROBERTS-RAPP, LISA
RUSSELL, JOHN C.
STROUPE, STEPHEN D.
: Abbott Laboratories
100 Abbott Park Road
                                                                                                                                                                                       COHEN, MAURICE
COLPITTS, TRACEY L.
FRIEDMAN, PAULA N.
GORDON, JULIAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8 amino acids
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                                                                                                                                                                              GRANADOS,
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                                         REAGENTS AND METHODS USEFUL FOR DETECTING DISEASES OF T
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Pred. No. 1.9e+06;
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Pred. No. 1.9e+06;
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RESULT 24
US-09-215-652-47
; Sequence 47, Application US/09215652
parent No. US20020045165A1
; OTHER INFORMATION: Affinity purification US-09-215-652-47
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APPLICANT: Edward N. Granados
APPLICANT: Steven C. Hodges
APPLICANT: Michael R. Klass
APPLICANT: Jon D. Kratochvil
APPLICANT: John C. Russell
APPLICANT: John C. Russell
APPLICANT: Stephen D. Stroppe
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Best Local S
                                                                                                           SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 47
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                                                                                                                                      TITLE OF INVENTION: Reagents and Methods Useful TITLE OF INVENTION: Breast FILE REPERROE: 6192 US. P1 CURRENT APPLICATION NUMBER: US/09/215,652 CURRENT FILING DATE: 1998-12-16 EARLIER FILING DATE: 1997-12-26 EARLIER FILING DATE: 1997-12-26 EARLIER FILING DATE: 1997-12-26 EARLIER FILING DATE: 1997-12-26
                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Tracey L. Colpitt APPLICANT: Paula N. Friedman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Maurice Cohen APPLICANT: Tracey L. Colp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Patricia Billing-Medel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
                                                                          TYPE: PR
                                              ORGANISM: Artificial Sequence
                                   FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Becker, Cheryl L.
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 61
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
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ATTORNEY/AGENT INFORMATION:
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OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
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TYPE: amino acid
STRANDEDNESS: sir
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OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
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Similarity 100.0%; Pred. No.
8; Conservative 0; Mismatch
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US-09-970-308-1
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                                                                    ; OTHER INFORMATION: Description of Artificial Sequence:Synthetic US-09-309-668A-1
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APPLICANT: BRIZZARD, BII
APPLICANT: HERNAN, RON
                                                                                                                                                      PRIOR APPLICATION NUMBER: 60/088,767
PRIOR FILING DATE: 1998-06-10
NUMBER OF SEQ ID NOS: 1
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 1
LENGTH: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1, Application US/09309668A Patent No. US20020045271A1
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Query Match
Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                   APPLICANT: Hussain, M. Mahmood
APPLICANT: Bakillah, Ahmed
TITLE OF INVENTION: Compounds and Methods for Identifying Compounds that
TITLE OF INVENTION: Interact with Microsomal Triglyceride Transfer Prote
TITLE OF INVENTION: Binding Sites on Apolipoprotein B and Modulate Lipit
TITLE OF INVENTION: Biosynthesis
FILE REFERENCE: MCP-0006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR FILING DATE: 1999-10-08
NUMBER OF SEQ ID NOS: 14
SOFTWARE: PatentIn Ver. 2.1
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CURRENT FILING DATE: 2001-10-03
PRIOR APPLICATION NUMBER: 09/415,000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: HERNAN, RON
TITLE OF INVENTION: PURIFICATION OF RECOMBINANT PROTEINS FUSED TO MULTIPLE
TITLE OF INVENTION: EPITOPES
FILE REFERENCE: SGM 6933.2
                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/309,668A CURRENT FILING DATE: 1999-05-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
                                                                                                           TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                       FEATURE:
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8; Conserv
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100.0%; ilarity 100.0%; Conservative 0
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Pred. No. 1.9e+06;
Score 47; DB 3;
Pred. No. 1.9e+06;
; Mismatches 0;
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Pred. No. 1.9e+06;
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                               Length 8;
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RESULT 28
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                                                                                                                                    US-09-817-413-1
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CURRENT PILLY APPLICATION NUMBER: US/09/855,722
CURRENT FILLY ADTE: 2001-05-16
PRIOR APPLICATION NUMBER: 09/214,278
PRIOR FILING DATE: 1999-01-26
NUMBER OF SEQ ID NOS: 32
SOFTWARE: PACENTIN Ver. 2.1
SEQ ID NO 25
                                                                            Query Match
Best Local S
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Patent No. US20020049306A1
GENERAL INFORMATION:
APPLICANT: Sakano, Seiji
APPLICANT: Itoh, Akira
                                                                                                                                                                                                                              PRIOR FILING DATE: 1999-04-03
NUMBER OF SEQ ID NOS: 25
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1, Application US/09817413 Publication No. US20020052478A1 GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                Matches
                                                                                                                                                                                                                                                                                                 APPLICANT: SRINIVISAN, ALGARSAMY
APPLICANT: KOPROWSK! HILARY
TITLE OF INVENTION: CHIMERIC VIRAL PROTEINS
FILE REFERENCE: Chimeric Viral Proteins
CURRENT APPLICATION NUMBER: US/09/817,413
CURRENT FILLING DATE: 2001-03-26
PRIOR APPLICATION NUMBER: 60/043,380
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                                                                                                                                                                            TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Description of Artificial Sequence: synthetic amino acid
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                                                                                                                                                OTHER INFORMATION: Description of Artificial Sequence:
                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                               LENGTH: 8
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                 1 DYKDDDDK 8
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                                                                100.0%; ilarity 100.0%; Conservative
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                                                                Score 47; DB 3;
Pred. No. 1.9e+06;
); Mismatches 0;
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Pred. No. 1.9e+06;
); Mismatches 0;
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RESULT 29 US-09-193-663-9

; Sequence 9, Application US/09193663

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RESULT 31
US-09-904-245-3
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                                                                                                                                                                       Sequence 3, Application US/09904245
Patent No. US20020076409A1
GENERAL INFORMATION:
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Patent No. US20020058310A1
GENERAL INFORMATION:
APPLICANT: Immunex Corporation
APPLICANT: Anderson, Dirk M
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Best Local (
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TITLE OF INVENTION: TNF-DELTA LIGAND AND USES THEREOF
FILE REFERENCE: 6255.US.02
CURRENT APPLICATION NUMBER: US/09/193,663
CURRENT FILING DATE: 1998-11-17
EARLIER APPLICATION NUMBER: 60/065,916
EARLIER FILING DATE: 1997-11-17
NUMBER OF SEQ ID NOS: 10
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 9
APPLICANT: March, Carl
APPLICANT: March, Carl
APPLICANT: Pluenneke, John
APPLICANT: O'Neal, Larry
TITLE OF INVENTION: METHOD FOR TREATING CANCER
FILE REFERENCE: 2943-A
CURRENT APPLICATION NUMBER: US/09/904,245
CURRENT FILING DATE: 2001-07-11
PRIOR APPLICATION NUMBER: 60/217,888
PRIOR FILING DATE: 2000-07-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Anderson, Dirk M
TITLE OF INVENTION: LECTIN SS3939 DNA AND POLYPEPTIDES
FILE REFERENCE: 2883-US
CURRENT APPLICATION NUMBER: US/09/887,855
CURRENT FILING DATE: 2001-06-22
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin version 3.1
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ORGANISM: Artificial Sequence
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8; Conserv
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APPLICANT: Tian, Hui
APPLICANT: Schultz, Joshua
APPLICANT: Schultz, Joshua
APPLICANT: Shan, Bei
APPLICANT: Tularik Inc.
TITLE OF INVENTION: Sitosterolemia Susceptibility Gene (SSITITLE OF INVENTION: Sitosterolemia Susceptibility Gene (SSITITLE OF INVENTION: MINDER: US/09/837,992
CURRENT APPLICATION NUMBER: US/09/837,992
CURRENT APPLICATION NUMBER: US/01/98,465
PRIOR APPLICATION NUMBER: US/01/98,465
PRIOR FILING DATE: 2000-04-18
PRIOR FILING DATE: 2000-04-18
PRIOR FILING DATE: 2000-05-15
NUMBER OF SEQ ID NOS: 45
SOFTWARE: Patentin Ver. 2.1
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                                                                                                                                                                                                                                                                                                 Sequence 45, Application Upper Patent No. US20020081687A1
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SEQ ID NO 15
LENGTH: 8
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Best Local Similarity
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LENGTH: 8
TYPE: PRT
ORGANISM: Artificial Sequence
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APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Patent No.
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TITLE OF INVENTION:
FILE REFERENCE: D60
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SOFTWARE: PatentIn vers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: Amino acid sequence of the FLAG octapeptide
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8; Conserv
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                                                                                                                                                                                                                                                                                                                      Application US/09837992
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Dmitriev, Igor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Curiel, David T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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DN: Epitope in the HI Loop of the Fiber Knob
D6080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Score 47; DB 3; 100.0%; Pred. No. 1.9e+06;
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Pred. No. 1.9e+06;
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GENERAL INFORMATION:
APPLICANT: Sheppard, Paul O.
ITITLE OF INVENTION:
FILE REFERENCE: 00-47
CURRENT APPLICATION NUMBER: US/09/923,995
CURRENT FILING DATE: 2001-08-07
PRIOR APPLICATION NUMBER: US 60/223,164
PRIOR PILING DATE: 2000-08-07
NUMBER OF SEQ ID NOS: 7
SOFTWARE: FastSEQ for Windows Version 3.0
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US-09-971-856-7
; Sequence 7, Application US/09871856
; Patent No. US20020081720A1
; GENERAL INFORMATION:
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Best Local
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE: OTHER INFORMATION: FLAG peptide tag
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 8
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: Apple Power Macintosh
OPERATING SYSTEM: Apple Operating System 7.5.5
SOFTWARE: Microsoft Word for Power Macintosh 6.

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/871,856
                                                                                                                                                                                                                                                                   Galibert, Laurent
Maraskovsky, Eugene
TITLE OF INVENTION: Receptor Activator of NF-kappaB
                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                        APPLICANT: Anderson, Dirk M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 рукрараж в
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8; Conserv
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                                                                                                                                                                                                    ADDRESSEE: Immunex Corporation, STREET: 51 University Street
                                                                                                                                  COUNTRY: USA
ZIP: 98101
                                                                                                                                                                    STATE: WA
                                                                                                                                                                                  CITY: Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 47; DB 3; 100.0%; Pred. No. 1.9e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 47; DB 3;
Pred. No. 1.9e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                      Law Department
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 40, Application US/09841894
Publication No. US20020086301A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: BILL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 7:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/996,139
FILING DATE: <UNKNOWN>
APPLICATION NUMBER: USSN 08/813,509
FILING DATE: 07 MARCH 1997
APPLICATION NUMBER: USSN 08/772,330
FILING DATE: 23 DECEMBER 1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
TELEFAX: (206)233-0644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLONE: FLAG peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 7:
                                                                                               COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Abbott Laboratories
STREET: 100 Abbott Park Road
                                                                                                                                                                                                                                                                                                                                                                                          GRANADOS, EDWARD N.
HODGES, STEVEN C.
KLASS, MICHAEL R.
KRATOCHVIL, JON D.
ROBERTS-RAPP, LISA
TITLE OF INVENTION: REAGENTS AND METHODS USEFUL
FOR DETECTING DISEASES OF THE PROSTATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: peptide
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/071,710
                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 рукрараж 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    8
                                        APPLICATION NUMBER: US/09/841,894
FILING DATE: 25-Apr-2001
CLASSIFICATION: «Unknown»
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS: No. US20020081720A1 Relevant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 2851-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Perkins, Patricia Anne
REGISTRATION NUMBER: 34,693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear
                                                                                                                                                                                                                             ZIP: 60064-3500
                                                                                                                                                                                                                                                                                    CITY: Abbott Park
                                                                                                                                                                                                                                                  COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COLPITTS, TRACEY L. FRIEDMAN, PAULA N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BILLING-MEDEL, PATRICIA COHEN, MAURICE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GORDON, JULIAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 47; DB 3; 100.0%; Pred. No. 1.9e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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US-09-991-681-32
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 32, Application US/099
Publication No. US20020086316A1
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 40:
SEQUENCE CHARACTERISTICS:
TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 32.
                                NAME: Becker, Cheryl L.
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 6084
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
TELEFAX: 847/938-2623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        / Match 100.0%; Score 47; DB 3; 1
Local Similarity 100.0%; Pred. No. 1.9e+06;
1es 8; Conservative 0; Mismatches 0;
                                                                                                                                                  APPLICATION NUMBER: 09/065,383
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                       OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/991,681
                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Abbott Laboratories
STREET: 100 Abbott Park Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KRATOCHVIL, JON D.

ROBERTS-RAPP, LISA
TITLE OF INVENTION: REAGENTS AND METHODS USEFUL
                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Abbott Park
STATE: IL
                                                                                                                                                                                                                                    FILING DATE: 26-Nov-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                         COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 60064-3500
                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 8 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Becker, Cheryl L. REGISTRATION NUMBER: 35,441
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: 847/935-1729
TELEFAX: 847/938-2623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/09991681
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BILLING-MEDEL, PATRICIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GRANADOS, EDWARD N. HODGES, STEVEN C. KLASS, MICHAEL R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COLPITTS, TRACEY L. FRIEDMAN, PAULA N. GORDON, JULIAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COHEN, MAURICE
                                                                                                                                                                                                                                                                                                                                                     IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FOR DETECTING DISEASES OF THE PROSTATE
                                                                                              6084.US.P1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6083.US.P1
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US-09-065-902-15
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                                                                             Matches
                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 15, Application US/09065902 Publication No. US20020086444A1 GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                                        TELEPHONE: (202) 371-260
TELEPAX: (202) 371-2540
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/044,262
FILING DATE: 24-APR-1997
ATTORNEY/AGENT INFORMATION:
NAME: Goldstein, Jorge A.
REGISTRATION NUMBER: 29,021
                                                                                                                                                                                         STRANDEDNESS: single TOPOLOGY: not relevant MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 0609.4270001/JAG/S-S TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 20005-3934
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein & Fox P.L.L.C.
STREET: 1100 New York Avenue, NW, Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: A Purified 20 kDa Presenilin 2
C-terminal Fragment and Methods of Screening for Compounds
that Inhibit Proteolysis of Presenilin 2
NUMBER OF SEQUENCES: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Tanzi,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE DESCRIPTION: SEQ ID NO: 32:
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                                                                                                                                                                                                                                                   TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/09/065,902 FILING DATE: 24-Apr-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                        LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acids
                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                       100.0%; Score 47; DB 3;
100.0%; Pred. No. 1.9e+06;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rudolph E.
                                                                                                                                                                                                                                                                                                                                                   371-2600
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Pred. No. 1.9e+06;
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                                                                                                             Length 8;
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                                                                       Indels
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                                                                                                                                                                                                         US-09-871-291-7
                                                                                                                                                                                                                                 RESULT 40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-865-363-7
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 7, Application US/09865363 Publication No. US20020086826A1
                                                                                                                                          Sequence 7, Application US/09871291
Publication No. US20020086827A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (206)233-06
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: Apple Power Macintosh

COMPUTER: Apple Power Macintosh

OPERATING SYSTEM: Apple Operating System 7.5.5

SOFTWARE: Microsoft Word for Power Macintosh 6.0.:

CURRENT APPLICATION DATA:

APPLICATION UNDBER: 20/0/865,363

FILING DATE: 25-May-2001

CLASSIFICATION: <Unknown>

PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: USSN 08/772,330
PILING DATE: 23 DECEMBER 1996
ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia Anne
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2852-A
TELECHMUNICATION INFORMATION:
1ELEPHONE: (206)587-0430
TELEPHONE: (206)233-0644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IMMEDIATE SOURCE:

CLONE: FLAG_ peptide
SEQUENCE DESCRIPTION: SEQ ID NO:
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TITLE OF INVENTION: Ligand for Receptor Activator of NF-kappaB
NUMBER OF SEQUENCES: 19
                                      Galibert, Laurent
Maraskovsky, Eugene
TITLE OF INVENTION: Receptor Activator of NF-kappaB
NUMBER OF SEQUENCES: 19
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                                                                                                                       APPLICANT: Anderson, Dirk M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE:
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APPLICATION NUMBER: USSN 08/813,509
FILING DATE: 07 MARCH 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: Not Relevant
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                                                                                                                                                                                                                                                                                                                                                                              Conservative
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Immunex Corporation, Law Department
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Pred. No. 1.9e+0;
Mismatches
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US-09-193-881-28
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                  SOFTWARE: FA
SEQ ID NO 28
LENGTH: 8
TYPE: PRT
                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Patricia Billing-Medel
APPLICANT: Maurice Cohen
APPLICANT: Tracey L. Colpitts
APPLICANT: Paula N. Friedman
                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 28, App. Publication No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
                                                                                                   APPLICANT: Edward N. Granados
APPLICANT: Michael R. Klass
APPLICANT: John C. Russell
APPLICANT: John C. Russell
APPLICANT: Stephen D. Stroupe
TITLE OF INVENTION: Reagents and Methods Useful for Detecting Diseases
TITLE OF INVENTION: Breast
FILE REFERENCE: 6248.US.Pl
CURRENT APPLICATION NUMBER: US/09/193,881A
CURRENT FILING DATE: 1998-11-18
NUMBER OF SEQ ID NOS: 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (206)233-0644
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
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APPLICATION NUMBER: US/09/871,291
FILING DATE: 30-May-2001
CLASSIFICATION: -Unknown>
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 98101

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Power Macintosh
OPERATING SYSTEM: Apple Operating System 7.
SOFTWARE: Microsoft Word for Power Macintos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE; amino acid
STRANDEDNESS: not relevant
TOPOLOGY: linear
MOLECULE TYPE: peptide
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION: TELEPHONE: (206)587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: USSN 08/772,330
FILING DATE: 23 DECEMBER 1996
ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia Anne
REGISTRATION NUMBER: 34,693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLONE: PLAG peptide SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 DYKDDDDK 8
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                                                                                     FastSEQ
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FILING DATE: <Unknown>
APPLICATION NUMBER: USSN 08/813,509
FILING DATE: 07 MARCH 1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 47; DB 3; 1 ilarity 100.0%; Pred. No. 1.9e+06; Conservative 0; Mismatches 0;
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                                                                                       for Windows Version 3.0
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CURRENT APPLICATION NUMBER: US/09/875,338
CURRENT FILING DATE: 2001-06-06
PRIOR APPLICATION NUMBER: 60/272,107
PRIOR FILING DATE: 2001-02-28
PRIOR APPLICATION NUMBER: 60/209,811
PRIOR FILING DATE: 2000-06-06
NUMBER OF SEQ ID NOS: 94
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 94
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US-09-875-338-94
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                                                                                                                                                                                          Sequence 7, Application US/09903248
Patent No. US20020102263A1
GENERAL INFORMATION:
APPLICANT: Wands, Jack R.
APPLICANT: de la Monte, Suzanne M.
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Best Local Similarity lou.
Whiches 8; Conservative
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            SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 7
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Best Local
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                                                                                       APPLICANT: Carlson, Rolf I.
TITLE OF INVENTION: DIAGNOSIS AND TREATMENT OF MALIGNANT NEOPLASMS
FILE REFERENCE: 21486-032 DIV5
CURRENT APPLICATION NUMBER: US/09/903,248
CURRENT FILING DATE: 2001-07-11
                                               PRIOR APPLICATION NUMBER: 09/436,184
PRIOR FILING DATE: 1999-11-08
NUMBER OF SEQ ID NOS: 9
                                                                                                                                                                                 APPLICANT:
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APPLICANT: PEACH, ROBERT
TITLE OF INVENTION: B7-RELATED NUCLEIC ACIDS AND
TITLE OF INVENTION: IMMUNOMODULATION
FILE REFERENCE: 3053-4071US2
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APPLICANT: CHANG, H
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ENGTH:
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8; Conser
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Pred. No. 1.9e+06;
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US-09-978-339-3
Sequence 3, Application US/09978339
Fatent No. US20020103358A1
GENERAL INFORMATION:
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APPLICANT: M. David
APPLICANT: Powers, Scott
APPLICANT: Tularik Inc.
APPLICANT: Tularik Inc.
FITLE OF INVENTION: KCNB: A No. US200201026
FILE REFERENCE: 018781-004010US
CURRENT APPLICATION NUMBER: US/09/798;584
CURRENT FILING DATE: 2001-03-03
PRIOR APPLICATION NUMBER: US 60/186,951
PRIOR FILING DATE: 2000-03-03
NUMBER OF SEQ ID NOS: 18
SOFTWARE: Patentin Ver. 2.1
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US-09-798-584-17
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Best Local Similarity
Marches 8; Conserva
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ENGTH: 8
                           COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple 7.1
SOFTWARE: Microsoft Word, Version
CURRENT APPLICATION DATA:
                                                                                                 ZIP: 98101
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                            Reddy, Pranhitha
TITLE OF INVENTION: No. US20020103358Alel Cytokine Designated Lerk-5
NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                APPLICANT: Cerretti, Douglas P.
                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
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8; Conserv
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                                                                                                                                                       STATE: Washington COUNTRY: US
                                                                                                                                                                                                         ADDRESSEE: Immunex Corporation STREET: 51 University Street
APPLICATION NUMBER: US/09/978,339 FILING DATE: 15-Oct-2001
                                                                                                                                                                                        CITY: Seattle
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Pred. No.
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PRIOR APPLICATION DATA:

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RESULT 47

US-09-903-063-7

; Sequence 7, Application US/09903063

; Patent No. US20020114810A1

; GENERAL INFORMATION:
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Best Local Similarity
Marches 8; Conserva
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US-09-859-604-7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Wands, Jack R.
APPLICANT: Wands, Jack R.
APPLICANT: Deutch, Alan H
APPLICANT: Deutch, Alan H
APPLICANT: Chanbari, Hossein A
TITLE OF INVENTION: DIAGNOSIS AND TREATMENT OF MALIGNANT NEOPLASMS
FILE REPERENCE: 21486-032 CIP
CURRENT APPLICATION NUMBER: US/09/859,604
CURRENT FILING DATE: 2001-05-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 7, Application US/09859604 Patent No. US20020110559A1
                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 09/436,184
PRIOR FILING DATE: 1999-11-08
NUMBER OF SEQ ID NOS: 13
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INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: Description of Artificial Sequence: FLAG epitope
                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 8
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IMMEDIATE SOURCE:
CLONE: FLAG peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Seese, Kathryn A.
REGISTRATION NUMBER: 32,172
REFERENCE/DOCKET NUMBER: 2823
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 587-0430
TELEFAX: (206) 233-0644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 08/271,948 FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                         1 DYKDDDDK 8
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STRANDEDNESS: single
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APPLICANT: Wands, Jack R.

APPLICANT: de la Monte, Suzanne M.

APPLICANT: Ince, Medim

APPLICANT: Carlson, Rolf I.

FITTLE OF INVENTION: DIAGNOSIS AND TREATMENT OF MALIGNANT NEOPLASMS

FILE REFERENCE: 21486-032 DIV2

CURRENT APPLICATION NUMBER: US/09/903,216

CURRENT FILING DATE: 2001-07-11

PRIOR APPLICATION NUMBER: 09/436,184

PRIOR APPLICATION NUMBER: 09/436,184

PRIOR FILING DATE: 1999-11-08

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 7

LENGTH: 8
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US-09-903-216-7
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                                                                                                                                                                                                                                                     B
                                                                                                                                                                   US-09-844-508-14
                                                                                                                                                                                       RESULT 49
                                                                                                                                                                                                                                                                                                                                                                                                                  , OTHER INFORMATION: Description of Artificial Sequence: FLAG US-09-903-216-7
Sequence 14, Application US/09844508
Patent No. US20020115215A1
GENERAL INFORMATION:
APPLICANT: WOLFFE, Alan P.
APPLICANT: COLLINGWOOD, Trevor
APPLICANT: COLLINGWOOD, Trevor
TITLE OF INVENTION: TARGETED MODIFICATION OF CHROMATIN STRUCTURE
FILE REFERENCE: 8325-0014 / S14-US1
CURRENT APPLICATION NUMBER: US/09/844,508
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Best Local (
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Best Local Similarity
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APPLICANT: Ince, Nedim
APPLICANT: Ince, Nedim
APPLICANT: Ince, Nedim
APPLICANT: Carlson, Rolf I.

TITLE OF INVENTION: DIAGNOSIS AND TREATMENT OF MALIGNANT NEOPLASMS
FILE REFERENCE: 21486-032 DIV3
CURRENT EPPLICATION NUMBER: US/09/903,063
CURRENT FILING DATE: 2001-10-11
RIOR APPLICATION NUMBER: 09/438,184
PRIOR FILING DATE: 1999-11-08
NUMBER OF SEQ ID NOS: 9
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TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
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8; Conserv
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Pred. No. 1.9e+06;
); Mismatches 0;
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Pred. No. 1.9e+06;
); Mismatches 0;
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                                                                                                                                                                                                                                             RESULT 51
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US-09-888-358-17
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             Sequence 7, Application US/09903199
Patent No. US20020122802A1
GENERRAL INFORMATION:
APPLICANT: Wands, Jack R.
APPLICANT: de la Monte, Suzanne M.
APPLICANT: Ince, Nedim
APPLICANT: Carlson, Rolf I.
TITLE OF INVENTION: DIAGNOSIS AND TREATMENT OF MALIGNANT NEOPLASMS
FILE REFERENCE: 21486-032 DIV4
CURRENT APPLICATION NUMBER: US/09/903,199
CURRENT FILING DATE: 2001-07-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: LEWIN, DAVID
APPLICANT: ADAMS, SEAN H.
APPLICANT: YU, XING XIAN
TITLE OF INVENTION: CGI-69 COMPOSITIONS AND METHODS OF
FILE REFERENCE: 10716/66
FILE REFERENCE: 10716/66
                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 8; Conserv
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Patent No. US20020119137A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/888,358
CURRENT FILING DATE: 2001-06-22
PRIOR APPLICATION NUMBER: 60/213,307
PRIOR FILING DATE: 2000-06-22
NUMBER OF SEQ ID NOS: 18
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PRIOR FILING DATE: 2000-08-28
NUMBER OF SEQ ID NOS: 49
SOFTWARE: Patentin Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Description of Artificial Sequence: FLAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: double-stranded oligonucleotide encoding OTHER INFORMATION: epitope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH:
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NUMBER: 09/436,184
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Pred. No. 1.9e+06;
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APPLICANT: SCHAMBYE, HANS T.
APPLICANT: VAN DEN HAZEL, BART
APPLICANT: VAN DEN HAZEL, BART
APPLICANT: CHRISTIANSEN, JESPER
APPLICANTION UNDRER: STIMULATING HORMONES
FILE REFERENCE: 31-00100US
CURRENT APPLICATION NUMBER: US/09/780,933
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: PA 2000 00220
PRIOR APPLICATION NUMBER: 60/104/035
PRIOR FILING DATE: 2000-02-22
PRIOR FILING DATE: 2000-02-22
PRIOR FILING DATE: 2000-02-22
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                                                                                                                                                                                                                                                                                                                                                                         US-09-780-933-20
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US-09-745-605-43
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US-09-903-199-7
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Matches
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                                                                                                                                                                                                                                                                                                                             Sequence 20, Application UPatent No. US20020127652A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 43, Application US/09745605 Patent No. US20020123617A1
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SEQ ID NO 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Starling, Gary C.
APPLICANT: Finger, Joshua N.
TITLE OF INVENTION: NOVEL IMMUNOGLOBIN SUPERFAMILY MEMBERS APEX-1, APEX-2,
TITLE OF INVENTION: AND APEX-3 AND USES THEREOF
FILE REFERENCE: DB13UM CURRENT SPLICATION NUMBER: US/09/745,605
CURRENT FILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: 60/172,025
PRIOR APPLICATION NUMBER: 60/172,025
PRIOR PILING DATE: 1999-12-23
NUMBER OF SEQ ID NOS: 44
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PRIOR APPLICATION NUMBER: PA 2000 01092
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TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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TYPE: PRT
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nes 8; Conserv
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ilarity 100.0%;
Conservative 0;
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Pred. No.
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CURRENT APPLICATION NUMBER: US/09/873,106B
CURRENT FILING DATE: 2001-06-01
PRIOR APPLICATION NUMBER: US 60/111,007
PRIOR FILING DATE: 1998-12-04
PRIOR APPLICATION NUMBER: US 60/115,647
PRIOR FILING DATE: 1999-01-13
PRIOR APPLICATION NUMBER: PCT/US99/26993
PRIOR APPLICATION NUMBER: PCT/US99/26993
PRIOR APPLICATION NUMBER: PCT/US99/26993
PRIOR FILING DATE: 1999-11-15
NUMBER OF SEQ ID NOS: 25
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 14
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; OTHER INFORMATION: Flag Epitope
US-09-873-106B-14
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US-09-995-593A-13
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PRIOR APPLICATION NUMBER: 60/225,
PRIOR FILING DATE: 2000-08-16
NUMBER OF SEQ ID NOS: 30
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 20
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                                                                                        Sequence 13, Application US/09995593A Patent No. US20020128197A1 GENERAL INFORMATION:
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Best Local
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TITLE OF INVENTION: Cloning and Characterization of a
TITLE OF INVENTION: Binding Protein (CD2BP2)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Reinherz, Ellis L. APPLICANT: Freund, Christian
APPLICANT: SAKANO, SEIJI
APPLICANT: ITOH, AKIRA
TITLE OF INVENTION: DIFFERENTIATION-SUPPRESSIVE POLYPEPTIDE
FILE REFERENCE: KP8447DIV
CURRENT APPLICATION NUMBER: US/09/995,593A
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TYPE: PRT
CORGANISM: Artificial Sequence
FEATURE:
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Similarity 100.0%;
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Nishizawa, Kazuhisa
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Pred. No.
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Pred. No.
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PRIOR APPLICATION NUMBER: 09/068,740
PRIOR TILING DATE: 1998-06-18
PRIOR APPLICATION NUMBER: JP 7-299611
PRIOR APPLICATION NUMBER: JP 7-299611
PRIOR APPLICATION NUMBER: JP 7-311811
PRIOR FILING DATE: 1995-11-30
PRIOR FILING DATE: 1995-11-30
PRIOR APPLICATION NUMBER: DCT/JP96/03356
PRIOR APPLICATION NUMBER: DCT/JP96/03356
PRIOR APPLICATION NUMBER: DCT/JP96/03356
PRIOR PILING DATE: 1996-11-15
PRIOR FILING DATE: 1996-11-15
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                                                                                                                                                                               RESULT 57
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US-09-900-530A-13
Sequence 37, Application US/09923246
Patent No. US20020128446A1
GENERAL INFORMATION:
APPLICANT: No. US20020128446A1ak, Julia
APPLICANT: Presnell, Scott R.
APPLICANT: Sprecher, Cindy A.
APPLICANT: Foster, Donald C.
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SEQ ID NO 13
LENGTH: 8
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/09/900,530A
CURRENT FILING DATE: 2002-03-19
PRIOR APPLICATION NUMBER: KR 2000-38441
PRIOR FILING DATE: 2000-07-06
NUMBER OF SEQ ID NOS: 48
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APPLICANT: Billiar, Timothy R.
APPLICANT: Billiar, Timothy R.
TITLE OF INVENTION: DNA Cassette for the Production of
TITLE OF INVENTION: Secretable Recombinant Trimeric Trail Proteins, Tetrae
TITLE OF INVENTION: /Doxycycline-Inducible Adeno-Associated Virus Vector,
TITLE OF INVENTION: Combination and Use in Gene Therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT ORGANISM: Artificial Sequence
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Pred. No. 1.9e+06;
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; FEATURE:
; OTHER INFORMATION: Synthetic
; FEATURE:
; NAME/KEY: misc feature
; OTHER INFORMATION: Synthetic
US-09-973-145-6
RESULT 59
US-09-825-561A-23
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"her 8; Conserv:
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                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn version 3.1
SEQ ID NO 6
LENGTH: 8
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Best Local Similarity
                                                                                                                                                                 Matches
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APPLICANT: Rothschild, Kenneth J.
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CURRENT APPLICATION NUMBER: US/09/973,145
CURRENT FILING DATE: 2001-10-09
PRIOR APPLICATION NUMBER: 09/382,950
PRIOR FILING DATE: 1999-08-25
PRIOR FILING DATE: 1999-08-25
NUMBER OF SEQ ID NOS: 18
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APPLICANT: Olejnik, Jerzy
TITLE OF INVENTION: N-Terminal and C-Terminal Markers in Nascent Proteins
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PRIOR APPLICATION NUMBER: US/09/923,246
CURRENT FILING DATE: 2001-08-03
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US/09/522,217
PRIOR FILING DATE: EARLIER FILING DATE: 2000-03-09
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US 60/123,904
PRIOR FILING DATE: EARLIER FILING DATE: 1999-03-11
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US 60/142,013
PRIOR FILING DATE: EARLIER FILING DATE: 1999-07-01
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SOFTWARE: FastSEQ for Windows Version 3.0
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APPLICANT: Dillon, Stacey R.
APPLICANT: Hammond, Angela K.
TITLE OF INVENTION: NOVEL CYTOKINE ZALPHA11 LIGAND
                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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                                                                                                                                                              Conservative
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100.0%; Pred. No.
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Pred. No.
                                                                                                                                                              Mismatches
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                                                                                                                                                                                                   Length 8;
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                                                                                                                                                            Indels
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US-09-825-561A-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 60/194,731
PRIOR FILING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: US 60/222,121
PRIOR FILING DATE: 2000-07-28
NUMBER OF SEQ ID NOS: 86
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: 00-22
CURRENT APPLICATION NUMBER: US/09/825,561A
CURRENT FILING DATE: 2000-04-05
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APPLICANT: No. US20020137677Alak,
APPLICANT: West, James W.
APPLICANT: Presnell, Scott R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Holly, Richard D.
APPLICANT: Nelson, Andrew J.
TITLE OF INVENTION: SOLUBLE ZALPHA11 CYTOKINE RECEPTORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                              ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
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                                                                                       SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS: Abbott Laboratories
ADDRESSEE: Abbott Laboratories
STREET: 100 Abbott Park Road
 PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
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Local Similarity 100.0%;
les 8; Conservativo
                                                         APPLICATION NUMBER: FILING DATE:
                                                                                                                            COMPUTER: IBM Compatible OPERATING SYSTEM: DOS
                                                                                                                                                                                                                   COUNTRY:
                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION:
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                                                                                                                                                                                                                       USA
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RUSSELL, JOHN C.
STROUPE, STEPHEN D.
VENTION: FEAGENTS AND METHODS USEFUL
VENTION: FOR DETECTING DISEASES OF THE GASTROINTESTINAL
VENTION: TRACT
                                                                                                                                                                                                                                                                                                                                                                                                                                                         HAYDEN, MARK
KLASS, MICHAEL R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BILLING-MEDEL, PATRICIA A. COHEN, MAURICE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COLPITTS, TRACEY L. FRIEDMAN, PAULA N.
08/828,845
                                                                      US/09/049,695A
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Pred. No. 1.9e+06;
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ATTORNEY/AGENT INFORMATION:

35,441

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Patent No. US20020142371A1
GENERAL INFORMATION:
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Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
NAME: Becker, Cheryl L.
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 6107
TELECOMPUNICATION INFORMATION:
TELEPHONE: 847/935-1729
TELEFAX: 847/938-2623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: RUSSELL, GAPPLICANT: STROUPE, SAPPLICANT: YU, HONG APPLICANT: YU, HONG TITLE OF INVENTION: H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT:
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APPLICANT:
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                                                                                                                                                                                                                          OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                             ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESSES: Abbott Laboratories
ADDRESSES: Abbott Park Road
STREET: 100 Abbott Park Road
                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/869,579
ETLING DATE: 05-JUN-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
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STRANDEDNESS: sir
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Becker, Cheryl L. REGISTRATION NUMBER: 35, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                 CITY: Abbott Park
STATE: IL
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ROBERTS-RAPP, LISA
RUSSELL, JOHN C.
STROUPE, STEPHEN D.
                                                                                                                                                                                                                                                                                                                                                   USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GRANADOS, EDWARD N. HODGES, STEVEN C. KLASS, MICHAEL R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GORDON,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BILLING-MEDEL, PATRICIA
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FOR DETECTING DISEASES OF THE URINARY TRACT
22
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                                                                                                                                                                                                          US/09/092,297
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                                                     6107.US.P1
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APPLICANT: Lyman, Stewart
APPLICANT: Armstrong, Allison
APPLICANT: Armstrong, Allison
APPLICANT: McKenna, Hilary
FILE REFERENCE: 03260.0087-00304/2881-WO
CURRENT APPLICATION NUMBER: US/09/852,391
CURRENT FILING DATE: 2001-05-09
PRIOR APPLICATION NUMBER: 60/108,452
PRIOR APPLICATION NUMBER: 60/108,452
INTOSER OF SEQ ID NOS: 5
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 62
US-09-903-023-7
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SEQ ID NO 7
LENGTH: 8
TYPE: PRT
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                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Sims, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 7, Application US/09903023 Patent No. US20020146421A1
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                                                                                                                                                                                                                                                                                                                            Sequence 3, Application US/09852391
Patent No. US20020146819A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 09/436,184
PRIOR FILING DATE: 1999-11-08
NUMBER OF SEQ ID NOS: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: 21486-032 DIV1
CURRENT APPLICATION NUMBER: US/09/903,023
CURRENT FILING DATE: 2001-10-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Wands, Jack R. APPLICANT: de la Monte,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Carlson, Rolf I.
TITLE OF INVENTION: DIAGNOSIS AND TREATMENT OF MALIGNANT NEOPLASMS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE: OTHER INFORMATION: Description of Artificial Sequence: FLAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
               ORGANISM: Artificial Sequence
                                  TYPE: PRT
FEATURE:
                                                   LENGTH: 8
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STRANDEDNESS: siz
TOPOLOGY: linear
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RESULT 65
US-09-263-959-985
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; ORGANISM: Homo sapiens
US-09-931-087A-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 64
US-09-931-087A-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 8; Conserv
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SEQ ID NO 19
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Best Local Similarity 100.0%;
Matches 8; Conservative 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 60/082,767
PRIOR FILING DATE: 1998-04-23
PRIOR APPLICATION NUMBER: US 60/113,296
PRIOR FILING DATE: 1998-12-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/931,087A
CURRENT FILING DATE: 2001-08-15
PRIOR APPLICATION NUMBER: 09/367,206
PRIOR FILING DATE: 1999-08-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Genentech, Inc.
TITLE OF INVENTION: NSP Molecules
FILE REFERENCE: P1223R1E
                                  ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/263,959
FILING DATE: 05-MAR-1999
                                                                                                                                                                                                                                                                                                                       APPLICANT: Hood, Leroy E.
APPLICANT: Rowen, Lee
APPLICANT: Koop, Ben F.
TITLE OF INVENTION: DIAGNOSTIC AND
NUMBER OF SEQUENCES: 1279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH:
                                                                                                                                                                                                                                                                                      ADDRESSEE: Seed and Berry LLP
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                      STREET:
                      CLASSIFICATION
                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                        CITY: Seattle
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                                                                                                                                                                                                                                                                    6300 Columbia Center, 701 Fifth Avenue
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ilarity 100.0%;
Conservative 0,
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Pred. No. 1.9e+06;
Mismatches 0;
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US-09-969-192-2
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US-09-969-192-2
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Best Local Similarity
Matches 8; Conserv
                                     Query Match
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                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 213564
INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 92
TELECOMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
TELEFAX: (206) 682-6031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/99/969,192
FILING DATE: 01-Oct-2001
PRIOR APPLICATION DATA:

'APPLICATION NUMBER: US 9-455061
FILING DATE: 06-DEC-1999
APPLICATION NUMBER: US 9-130225
FILING DATE: 06-AUG-1998
APPLICATION NUMBER: US 8-701124
FILING DATE: 21-AUG-1996
ATTORNEY/AGENT INFORMATION:

NAME: Hefner, M. Daniel
REGISTRATION NUMBER: 41,826
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
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                                                                                            MOLECULE TYPE: peptide SEQUENCE DESCRIPTION:
                                                                                                               TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 60601
COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Leydig, Voit & Mayer, Ltd.
STREET: Two Prudential Plaza - 49th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: WICKHAM, THOMAS J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 DYKDDDDK 8
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Similarity 100.0%;
8; Conservative 0
                                                                                                                                                TYPE: amino acid
STRANDEDNESS: single
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
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STATE: Illinois
100.0%;
ilarity 100.0%;
Conservative 0
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                 Score 47; DB 3; 1
Pred. No. 1.9e+06;
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Pred. No. 1.9e+06;
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                                     Length 8;
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US-09-843-245-12
; Sequence 12, Application US/09843245
; Patent No. US20020164672A1
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                                                           RESULT 69
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LENGTH: 8
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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APPLICANT: Bein, Kiflai
APPLICANT: Simons, Mic
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Patent No. US20020159989A1
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SEQ ID NO 20
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Matches 8; Conser
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Best Local &
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APPLICANT: MACH, Bernard
TITLE OF INVENTION: New Transcription Factor of MHC Class II Genes, Substance
TITLE OF INVENTION: Capable of Inhibiting This New Transcription Factor and
TITLE OF INVENTION: Medical Uses of These Substances
                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Simons, Michael
TITLE OF INVENTION: Peptide Inhibitor of MMP Activity and Angiogenesis
FILE REFERENCE: BIT-003
CURRENT APPLICATION NUMBER: US/09/766,700A
CURRENT FILING DATE: 2001-01-22
NUMBER OF SEQ ID NOS: 8
SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: 010830-117
CURRENT APPLICATION NUMBER: US/09/840,243B
CURRENT FILING DATE: 2001-04-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: EP 98120085.0 PRIOR FILING DATE: 1998-10-24 NUMBER OF SEQ ID NOS: 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE: OTHER INFORMATION: Description of Artificial Sequence:FLAG epitope
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                                                                                                                                         1 DYKDDDDK 8
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Pred. No. 1.9e+06;
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TITLE OF INVENTION: No. US20020164716Alel TNE
TITLE OF INVENTION: inhibitors of ligand bir
FILE REFERENCE: GFN-523CP4DV3
CURRENT APPLICATION NUMBER: US/09/989,350
CURRENT FILING DATE: 2001-11-20
PRIOR APPLICATION NUMBER: 09/185,258
PRIOR FILING DATE: 1998-11-02
PRIOR FILING DATE: 1998-11-02
PRIOR APPLICATION NUMBER: 08/839,032
PRIOR APPLICATION NUMBER: 08/698,551
PRIOR APPLICATION NUMBER: 08/698,551
PRIOR APPLICATION NUMBER: 08/602,228
PRIOR FILING DATE: 1996-08-15
PRIOR FILING DATE: 1996-08-15
PRIOR APPLICATION NUMBER: 08/602,28
PRIOR FILING DATE: 1996-08-15
PRIOR APPLICATION NUMBER: 08/602,28
PRIOR APPLICATION NUMBER: 08/602,28
PRIOR APPLICATION NUMBER: 08/602,28
PRIOR APPLICATION NUMBER: 08/602,28
PRIOR APPLICATION NUMBER: 08/603,751
PRIOR APPLICATION NUMBER: 08/604,440
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-19
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US-09-989-350-19
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                                                                                                                                                                               ; OTHER INFORMATION: Description of Artificial Sequence: Flag-tag US-09-989-350-19
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Patent No. US20020164716A1
GENERAL INFORMATION:
APPLICANT: Lin, Lih-Ling
APPLICANT: Chen, Jennife
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 12
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Best Local
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CURRENT FILING DATE: 2001-04-25
NUMBER OF SEQ ID NOS: 13
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APPLICANT: Ramjaun, Authorie Rachid
TITLE OF INVENTION: REGULATION OF JNK ACTIVITY BY MODULATION OF THE
TITLE OF INVENTION: INTERACTION BETWEEN THE ENDOCYTIC PROTEIN ENDOPHILIN
TITLE OF INVENTION: AND THE GERMINAL CENTER KINASE-LIKE KINASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 08/327,514
PRIOR FILING DATE: 1994-10-19
NUMBER OF SEQ ID NOS: 19
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                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Ver.
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FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: FLAG
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                                                                                                                                                                                                                             ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                             TYPE: PRT
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ilarity 100.0%;
Conservative (
                                                                                           Conservative
                                                                                                              100.0%; Score 47;
100.0%; Pred. No.
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Pred. No. 1.9e+06;
); Mismatches 0;
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                                                                                                                 1.9e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-858-332-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 7, Application US/09877650 Patent No. US20020169117A1 GENERAL INFORMATION:
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Patent No. US20020164718A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/858,332
CURRENT FILING DATE: 2002-07-02
PRIOR APPLICATION NUMBER: 09/404,017
PRIOR FILING DATE: 1999-09-23
PRIOR APPLICATION NUMBER: 60/101,867
PRIOR FILING DATE: 1998-09-25
NUMBER OF SEQ ID NOS: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Tchaga, Grigory S.
APPLICANT: Jokhadze, George
TITLE OF INVENTION: Metal Ion Affinity Tags and Methods for
TITLE OF INVENTION: Using the Same
FILE REFERENCE: CLON056CIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: synthetic peptide
                             COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: Apple Power Macintosh

OPERATING SYSTEM: Apple Operating System 7.5.5

SOFTWARE: Microsoft Word for Power Macintosh 6.0.1

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/877,650

FILING DATE: 08-Jun-2001

CLASSIFICATION UMBER: 08/95,659

FILING DATE: 1997-12-22

APPLICATION NUMBER: USSN 08/813,509

FILING DATE: 1997-12-22

APPLICATION NUMBER: USSN 08/813,509

FILING DATE: 07 MARCH 1997

APPLICATION NUMBER: USSN 08/772,330

FILING DATE: 23 DECEMBER 1996

ATTORNEY/AGENT IMPORMATION:

NAMME: Dervices Decripts Anno
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nes 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Ligand for Receptor Activator of NF-kappaB NUMBER OF SEQUENCES: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Anderson, Dirk M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 рукрараж в
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
ZIP: 98101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: WA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Immunex Corporation, Law Department STREET: 51 University Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ДУКДДДДДК 8
NAME: Perkins, Patricia Anne REGISTRATION NUMBER: 34,693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%;
ilarity 100.0%;
Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Maraskovsky, Eugene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Galibert,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0,
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Pred. No. 1.9e+06;
Pred. No. 1.9e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 3; Length 8;
                                                                                                                                                                                                                                                                                                           Macintosh 6.0.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-981-636-1
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                                                                                                                                                                          Sequence 19, Application US/09092296 Publication No. US20020188114A1
                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 1
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                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: MARKS, JAMES
APPLICANT: MIELSEN, ULRIK
APPLICANT: KIRDOIN, DMITRI
TITLE OF INVENTION: METHODS OF HIGH-THROUGHPUT SCREENING FOR INTERNALIZING ANTIBODIE
FILE REFERENCE: 4077-897710US
CURRENT APPLICATION NUMBER: US/09/981,636
CURRENT FILING DATE: 2002-06-25
PRIOR APPLICATION NUMBER: US 60/241,279
PRIOR APPLICATION NUMBER: US 60/241,279
PRIOR FILING DATE: 2000-10-18
NUMBER OF SEQ ID NOS: 3
SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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                                                                                                                                                              GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH:
                                                                                                                            APPLICANT:
APPLICANT:
                                                              APPLICANT:
               TITLE OF INVENTION:
                                                                                              APPLICANT:
                                                                                                               APPLICANT:
NUMBER OF SEQUENCES:
                                                APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 2852-A TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                          1 DYKODDDK 8
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STRANDEDNESS: not relevant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (206) 587-04:
TELEFAX: (206) 233-0644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
                                                            COHEN, MAURICE
COLPITTS, TRACEY L.
FRIEDMAN, PAULA N.
KLASS, MICHAEL R.
RUSSELL, JOHN C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ENGTH: 8 amino acids
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                                        STROUPE, STEPHEN D.
                                                           RUSSELL,
                                                                                                                                           BILLING-MEDEL, PATRICIA
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REAGENTS AND METHODS USEFUL FOR DETECTING DISEASES OF THE LUNG 20
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Pred. No.
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Pred. No. 1.9e+06;
                                                                                                                                                                                                                                                                                                                                        Mismatches
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hes 0;
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Abbott Laboratories

ADDRESSEE: Abbott STREET: 100 Abbott CITY: Abbott Park

100 Abbott Park Road

COUNTRY:

USA

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RESULT 75
US-09-104-408-36
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Publication No. US20020188115A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEX:
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                        ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Billing-Medel, Patricia A. et al.
TITLE OF INVENTION: REAGENTS AND METHODS USEFUL
TITLE OF INVENTION: FOR DETECTING DISEASES OF THE BREAST
NUMBER OF SEQUENCES: 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Becker, Cheryl L.
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 6104.US.01
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/0
FILING DATE: 05-JUN-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 37
CORRESPONDENCE ADDRESS:
ADDRESSEE: Abbott Laboratories
STREET: 100 Abbott Park Road
               CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 6122.US.01
FILING DATE: 25-JUN-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: FILING DATE: CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 847/938-2623
                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                          CITY: Abbott Park
STATE: IL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH:
                                                                                                        APPLICATION NUMBER:
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Pred. No. 1.9e+06;
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RESULT 76
US-09-861-097-15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 100.0%; Score 47; DB 3; Best Local Similarity 100.0%; Pred. No. 1.9e+06; Matches 8; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS
                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 15: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Becker, Cheryl L
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/09/861,097
FILING DATE: 18-May-2001
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 09/220,602
FILING DATE: 25-MAR-1994
ATTORNEY/AGENT INFORMATION:
NAME: Haile, Ph.D., Lisa A.,
REGISTRATION NUMBER: 38,347
REFERENCE/DOCKET NUMBER: 07257/015001(PD3205)
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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STRANDEDNESS: sin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: FISH & RICHARDSON P.C.
STREET: 4225 Executive Square, Suite 1400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DERIJARD, BENOIT
TITLE OF INVENTION: ONCOPROTEIN PROTEIN KINASE
NUMBER OF SEQUENCES: 25
CORRECTORING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: KARIN, MICHAEL
SEQUENCE DESCRIPTION: SEQ ID NO: 15:
                                                       FEATURE:
                                                                           MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DYKDDDDK 8
                                                                                                                  TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                    TELEPHONE: (619) 678-5099
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: La Jolla
STATE: California
                                  NAME/KEY: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 92037
                                                                                                  TOPOLOGY:
                                                                                                                                                       LENGTH: 8 amino acids
                    LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8 amino acids
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TNG, LIN
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[KO, HIBI
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; OTHER INFORMATION: Description of Artificial Sequence: synthetic ; OTHER INFORMATION: peptide sequence US-09-908-943A-186
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APPLICANT: Klein, Elliott S.
APPLICANT: Chandraratna Roshantha A.
TITLE OF INVENTION: Methods of Detecting Dissociated Nuclear
TITLE OF INVENTION: Hormone Receptor Ligands
FILE REFERENCE: P-AR 4528
CURRENT APPLICATION NUMBER: US/09/814,604
CURRENT FILING DATE: 2001-03-22
NUMBER OF SEQ ID NOS: 52
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 5
  Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 8; Conservative 0
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                                                                                                                                                                                                                        SEQ ID NO 186
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Best Local
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APPLICANT: Emmons, Thomas L.
APPLICANT: Bienkowski, Mike J.
APPLICANT: Heinrikson, Robert L.
TITLE OF INVENTION: SUBSTRATES AND ASSAYS FOR BETA-SECRETASE ACTIVITY
FILE REFERENCE: 29915/00281A.US1
CURRENT APPLICATION NUMBER: US/09/908,943A
CURRENT FILING DATE: 2001-07-19
PRIOR APPLICATION NUMBER: 05/219,795
PRIOR APPLICATION NUMBER: 05/219,795
PRIOR FILING DATE: 2000-07-19
NUMBER OF SEQ ID NOS: 197
NUMBER OF SEQ ID NOS: 197
SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Yan, Riqiang APPLICANT: Tomasselli, Alfredo G.
                                                                                                                                           TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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les 8; Conservative
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Score 47; DB 3; Le
Pred. No. 1.9e+06;
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Pred. No. 1.9e+06;
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US-09-997-623-47
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US-09-997-623-47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Maxygen Aps; Maxygen Holdings
TITLE OF INVENTION: Protein C or activated protein C-like molecules
FILE REFERENCE: 02194410 - protein C
FULE REFERENCE: 02194410 - protein C
CURRENT APPLICATION NUMBER: US/09/997,623
CURRENT FILING DATE: 2001-11-29
CURRENT FILING DATE: 2001-11-29
                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 50
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 12
                                     Matches
                                                    Query Match
Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 12, Application US/09995898A Publication No. US20030027253A1
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Best Local
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Publication No. US20030018175A1
                                                                                                                                                                                                                                                                 FILE REFERENCE: 00-108
CURRENT APPLICATION NUMBER: US/09/995,898A
CURRENT FILING DATE: 2001-11-28
PRIOR APPLICATION NUMBER: US 60/253,561
PRIOR FILING DATE: 2000-11-28
PRIOR APPLICATION NUMBER: US 60/267,211
PRIOR FILING DATE: 2001-02-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Presnell, Scott R. APPLICANT: Xu, Wenfeng APPLICANT: No. US20030027253
                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Grant, Francis J.
TITLE OF INVENTION: CYTOKINE RECEPTOR ZCYTOR19
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PRIOR FILING DATE: 2001-10-17
NUMBER OF SEQ ID NOS: 48
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                                                                                                                                                                                LENGTH: 8
TYPE: PRT
                                                                                                                                             ORGANISM: Artificial Sequence FEATURE:
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TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                            OTHER INFORMATION: FLAG peptide tag
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                                                        Similarity
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8; Conservative
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No. US20030027253Alak, Julia E.
Whitmore, Theodore E.
                                     Conservative
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                                                      Score 47;
Pred. No.
                                     Mismatches
                                                        1.9e+06;
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RESULT 83
US-09-950-634-23
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Best Local Similarity
Marches 8; Conserve
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APPLICANT: Maxygen Aps; Maxygen Holdings
TITLE OF INVENTION: Protein C or activated protein C-like molecules
FILE REFERENCE: 0219us310 - protein C
CURRENT APPLICATION NUMBER: US/09/978,917A
CURRENT FILING DATE: 2001-10-17
NUMBER OF SEQ ID NOS: 48
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 47
                                                                                                Sequence 23, Application US/09950634 Publication No. US20030032775A1 GENERAL INFORMATION:
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Publication No. US20030031670A1
GENERAL INFORMATION:
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Publication No.
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SEQ ID NO 7
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Best Local Similarity
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TITLE OF INVENTION: DIAGNOSIS AND TREATMENT OF MALIGNANT NEOPLASMS
FILE REFERENCE: R.I. Hosp. - Malignant Neoplasms
CURRENT APPLICATION NUMBER: US/09/436,184
CURRENT FILING DATE: 1999-11-08
NUMBER OF SEQ ID NOS: 7
SOFTWARE: Patentin Ver. 2.0
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TYPE: PRT
ORGANISM: Artificial Sequence
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 TITLE OF
                                                                                  APPLICANT:
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de la Monte, Suzanne M.
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INVENTION: EFFECTOR PROTEINS OF RAPAMYCIN
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                                                                                Molnar-Kimber, Katherine
                                               Caggiano, Thomas J.
                            Nakanishi,
                                                                 Failli, Amedeo F.
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                             Koji
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Pred. No. 1.9e+06;
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Pred. No. 1.9e+06;
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                                                                                                                                                                                                                                                                   Sequence 225, Application US/09, Publication No. US20030044404A1 GENERAL INFORMATION:
APPLICANT: Rebar, Edward APPLICANT: Jamieson, Andrew APPLICANT: Liu, Qiang
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APPLICANT: Jarvis, Eric
APPLICANT: Sangamo BioSciences, Inc.
TITLE OF INVENTION: Regulation of Angiogenesis With Zinc
TITLE OF INVENTION: Finger Proteins
FILE REFERENCE: 019496-005820US
CURRENT APPLICATION NUMBER: US/09/846,033B
CURRENT FILING DATE: 2001-04-30
PRIOR APPLICATION NUMBER: US 09/733,604
PRIOR FILING DATE: 2000-12-07
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 08/471,112
FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/384,524
FILING DATE: 13-FEB-1995
APPLICATION NUMBER: US 08/312,023
FILING DATE: 26-SEP-1995
APPLICATION NUMBER: US 08/207,975
APPLICATION NUMBER: US 08/207,975
FILING DATE: 08-MAR-1994
ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear MOLECULE TYPE: peptide SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION: TELEPHONE: 202-408-4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/09/950,634
FILING DATE: 13-Sep-2001
CLASSIFICATION: cUnknown>
FRIOR APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 рукорорк в
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                                                                                                                                                                           Liu, Qiang
Liu, Pei-Qi
Wolffe, Alan
Eisenberg, Stephen I
Jarvis, Eric
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Washington
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STREET: 1300 I Street, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                              Application US/09846033B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 47; DB 3; ilarity 100.0%; Pred. No. 1.9e+06; Conservative 0; Mismatches 0;
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Publication No. US2003
GENERAL INFORMATION:
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 225
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                                                                                                                                                                                                                 TELEFAX: (619) 678-5099
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                    NAME: Haile, Ph.D., Lisa A.,
REGISTRATION NUMBER: 38,347
REFERENCE/DOCKET NUMBER: 07257/015001(PD3205)
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Patentin Release #1.0, V
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/861,098
FILING DATE: 18-May-2001
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 08/220,602
FILING DATE: 25-MAR-1994
ATTORNEY/AGENT INFORMATION:
                                                  SEQUENCE DESCRIPTION: SEQ ID
                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                           MOLECULE TYPE:
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TITLE OF INVENTION: ONCOPROTEIN PROTEIN KINASE
NUMBER OF SEQUENCES: 25
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                                                                                                                                          LENGTH: 8 amino acids TYPE: amino acid STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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STATE: California
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                                                                                     NAME/KEY: Peptide
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No. US20030044788A1
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ANNING, LIN
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25 Executive Square, Suite 1400
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Pred. No. 1.9e+06;
; Mismatches 0;
 Score 47;
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DB 3;
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Length
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APPLICANT: Tularik Inc.

APPLICANT: Board of Regents, The University of Texas System
ITILE OF INVENTION: ABCG5 and ABCG8: Compositions and Methods of
FILE REFERENCE: 018781-007320US
CURRENT APPLICATION NUMBER: US/09/989,981A
CURRENT FILING DATE: 2002-07-23
PRIOR APPLICATION NUMBER: US 60/252,235
PRIOR APPLICATION NUMBER: US 60/253,645
PRIOR FILING DATE: 2000-11-28
PRIOR FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.1
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Best Local Similarity
Thes 8; Conserve
                                      US-09-993-180-33
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US-09-989-981A-12
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                                                                                                              PRIOR APPLICATION NUMBER: US 60/282,745
PRIOR FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 46
SOFTWARE: PatentIn version 3.0
SEQ ID NO 33
                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
  Query Match
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Publication No.
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                                                                                                                                                                                                          CURRENT FILING DATE: 2001-11-14
PRIOR APPLICATION NUMBER: US 60/248,434
PRIOR FILING DATE: 2000-11-14
PRIOR APPLICATION NUMBER: US 60/257,610
PRIOR FILING DATE: 2000-12-21
                                                                                                                                                                                                                                                                                    APPLICANT: Bristol-Myers Squibb Company
TITLE OF INVENTION: POLYNUCLEOTIDE ENCODING A NOVEL HUMAN SERPIN SECRETED FROM LYMPH
TITLE OF INVENTION: CELLS, LSI-01
FILE REFERENCE: DO051.NP
CURRENT APPLICATION NUMBER: US/09/93,180
CURRENT FILING DATE: 2001-11-14
                                                                              LENGTH: 8
TYPE: PRT
                                                      ORGANISM: Bacteriophage T7
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100.0%; Pred. No.
tive 0; Mismatch
  100.0%;
  Score 47;
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DB
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Length 8;
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US-09-999-2208-9
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                                                                                                                                                          ORGANISM: bacteriophage T7 US-09-999-220B-9
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Publication No. US20030059923A1
GENERAL INFORMATION:
APPLICANT: Bristol-Myers Squibb Company
TITLE OF INVENTION: POLYNUCLEOTIDE ENCODING A NOVEL HUMAN POTASSIUM CHANNEL ALPHA-SUE
TITLE OF INVENTION: K+alpham1, AND VARIANTS THEREOF
FILE REFERENCE: D0050NP
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Matches 8; Conserv
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                                                                                                 Query Match
Best Local Similarity
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APPLICANT: Tularik Inc.
APPLICANT: Tularik Inc.
TITLE OF INVENTION: IRAK-4: Compositions and Methods of Use
FILE REFERENCE: 018781-003910US
CURRENT APPLICATION NUMBER: US/09/759,595
CURRENT FILING DATE: 2001-01-13
PRIOR APPLICATION NUMBER: US 60/176,395
PRIOR FILING DATE: 2000-01-13
NUMBER OF SEQ ID NOS: 7
COMMENCE OF SEQ ID NOS: 7
                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/99,220B
CURRENT FILING DATE: 2002-06-04
PRIOR APPLICATION NUMBER: US 60/245,383
PRIOR FILING DATE: 2000-11-02
PRIOR APPLICATION NUMBER: US 60/257,780
PRIOR FILING DATE: 2000-12-21
PRIOR APPLICATION NUMBER: US 60/269,854
PRIOR APPLICATION NUMBER: US 60/269,854
PRIOR FILING DATE: 2001-02-20
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                                                                                                                                                                                                         LENGTH: 8
TYPE: PRT
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                          DYKDDDDK 8
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DYKDDDDK
                                                                       100.0%; Score 47; DB 3; illarity 100.0%; Pred. No. 1.9e+06; Conservative 0; Mismatches o.
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US-09-880-748-3238
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CURRENT APPLICATION NUMBER: US/09/880,748
CURRENT FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/212,210
PRIOR PILING DATE: 2000-06-15
PRIOR PILING DATE: 2000-06-15
PRIOR APPLICATION NUMBER: 60/240,816
PRIOR PILING DATE: 2000-10-17
PRIOR PILING DATE: 2001-03-16
PRIOR FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR PILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR PILING DATE: 2001-03-25
PRIOR PILING DATE: 2001-03-25
PRIOR PILING DATE: 2001-03-26
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Publication No. US20030059937A1
GENERAL INFORMATION:
APPLICANT: Ruben et al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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Publication No.
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Best Local Similarity
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APPLICANT: ANDERS
APPLICANT: HANSEN
                                                                    PRIOR APPLICATION NUMBER: DK PA 2000 00024
PRIOR FILING DATE: 2000-01-10
PRIOR APPLICATION NUMBER: DK PA 2000 00341
PRIOR FILING DATE: 2000-03-02
PRIOR APPLICATION NUMBER: DK PA 2000 00943
PRIOR APPLICATION NUMBER: DK PA 2000 00943
PRIOR FILING DATE: 2000-06-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: ANDERSEN, KIM VILBOUR APPLICANT: HANSEN, CHRISTIAN KARSTEN APPLICANT: MIKKELSEN, JAN MOLLER TITLE OF INVENTION: G-CSF CONJUGATES FILE REFERENCE: 31.000700US
                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/904,196B CURRENT FILING DATE: 2001-07-11 PRIOR APPLICATION NUMBER: US/09/760,008 PRIOR FILING DATE: 2001-05-11 PRIOR APPLICATION NUMBER: 60/176,376 PRIOR FILING DATE: 2000-01-14 PRIOR FILING DATE: 2000-01-14 PRIOR FILING DATE: 2000-01-14
NUMBER OF SEQ ID NOS: 15
SOFTWARE: PatentIn Ver. :
SEQ ID NO 14
                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/189,506
PRIOR FILING DATE: 2000-03-15
PRIOR APPLICATION NUMBER: 60/215,644
PRIOR FILING DATE: 2000-06-30
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TYPE: PRT
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NAME/KEY: site
LOCATION: (1)..(8)
OTHER INFORMATION: Flag Tag
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Pred. No. 1.9e+06
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Sequence 39, Application US/0992226

Publication No. US20030077664A1

GENERAL INFORMATION:
APPLICANT: Zhao, Yi
APPLICANT: Thacher, Scott M.
APPLICANT: Xiao, Jia-Hao
APPLICANT: Kiao, Jia-Hao
APPLICANT: Kusari, Jyotirmoy
APPLICANT: Kusari, Jyotirmoy
APPLICANT: Chandraratna, Roshantha A.
ITILE OF INVENTION: Methods of Screening For Compounds That
ITILE OF INVENTION: Modulate Hormone Receptor Activity
FILE REFERENCE: P-AR 4681
CURRENT APPLICATION NUMBER: US/09/922,226
CURRENT PILING DATE: 2002-01-09
PRIOR APPLICATION NUMBER: US 60/284,797
PRIOR APPLICATION STEEL 2002-04-18
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US-09-922-226-39
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CURRENT FILING DATE: 2001-10-04
PRIOR APPLICATION NUMBER: 09/263,022
PRIOR FILING DATE: 1999-03-05
PRIOR APPLICATION NUMBER: 08/843,704
PRIOR APPLICATION NUMBER: 08/843,704
PRIOR FILING DATE: 1997-04-16
PRIOR APPLICATION NUMBER: 08/842,898
PRIOR APPLICATION NUMBER: 08/842,898
PRIOR APPLICATION NUMBER: 08/842,898
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Best Local :
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: McCarthy, Sean A.
TITLE OF INVENTION: MOVEL HUMAN DICKKOPF-RELATED PROTEIN AND NUCLEIC ACID
TITLE OF INVENTION: MOLECULES AND USES THEREFOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Ver.
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PRIOR FILING DATE: 1998-01-15
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8; Conser
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Pred. No. 1.9e+06;
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Pred. No. 1.9e+06;
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PRIOR TILING DATE: 1999-09-30
PRIOR APPLICATION NUMBER: 09/01
PRIOR APPLICATION NUMBER: 09/01
PRIOR PILING DATE: 1998-01-22
PRIOR PILING DATE: 1997-10-06
PRIOR PILING DATE: 1998-01-27
PRIOR FILING DATE: 1998-01-27
PRIOR FILING DATE: 1998-01-27
PRIOR FILING DATE: 1997-10-06
PRIOR PILING DATE: 1997-10-06
PRIOR PILING DATE: 1997-10-06
PRIOR PILING DATE: 1997-10-06
PRIOR PILING DATE: 2000-12-21
PRIOR FILING DATE: 2000-12-21
PRIOR FILING DATE: 2000-12-21
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Best Local S
Matches 8
                                                                                      Query Match
Best Local (
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                                                                      Matches
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LENGTH: 8
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PRIOR FILING DATE: 1998-12-18
PRIOR APPLICATION NUMBER: 08/9
PRIOR FILING DATE: 1997-12-19
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TITLE OF INVENTION: NOVEL EDIRF, MTR-1, LSP-1,
TITLE OF INVENTION: AND USES THEREFOR
FILE REFERENCE: MNI-107CP2
CURRENT APPLICATION NUMBER: US/09/774,381
CURRENT FILING DATE: 2001-01-30
CURRENT FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                    LENGTH: 8
TYPE: PRT
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                                                                                                                                                         ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence:flag
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ORGANISM: Artificial
FEATURE:
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FILING DATE: 2000-01-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: FILING DATE: 1998-0
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FILING DATE: 1997-10-06
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                                                                                        Similarity
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8; Conserv
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                                                                      Conservative
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                                                                                        100.0%;
                                                                                    100.0%; Score 47; DB 3; 100.0%; Pred. No. 1.9e+06;
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US-09-876-790-11; Sequence 11, App.; Publication No.
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Best Local Similarity
Watches 8; Conserve
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US-09-876-790-11
RESULT 97
US-09-782-587B-18
; Sequence 18, Application US/09782587B
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PRIOR APPLICATION NUMBER: 60/112,163
PRIOR ETLING DATE: 1998-12-14
PRIOR APPLICATION NUMBER: 60/146,675
PRIOR APPLICATION NUMBER: PCT/US99/29549
PRIOR FILING DATE: 1999-12-14
NUMBER OF SEQ ID NOS: 15
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APPLICANT: Rosen, Craig A.

APPLICANT: Rosen, Craig A.

TITLE OF INVENTION: BINDING POLYPEPTIDES AND METHODS BASED THEREON
FILE REFERENCE: DYX-025.1 PCT; DYX-025.1 US

CURRENT APPLICATION NUMBER: US/09/932,613

CURRENT FILING DATE: 2001-08-17

NUMBER OF SEQ ID NOS: 458

SOFTWARE: Patentin version 3.1

SEQ ID NO 183
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Best Local
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Publication No. US20030091565A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: SIMS, John E.
APPLICANT: SMITH, Dirk E.
APPLICANT: BORN, Teresa L.
TITLE OF INVENTION: IL-1 ZETA, IL-1 ZETA;
FILE REFERENCE: 2008-US
CURRENT APPLICATION NUMBER: US/09/876,790
CURRENT FILING DATE: 2002-09-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Human Genome Sciences, APPLICANT: Beltzer, James P. APPLICANT: Potter, M. Daniel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT ORGANISM: Artificial sequence FEATURE:
                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                      TYPE: PRT
                                                                                                                                                                                                                                                              OTHER INFORMATION: epitope tag
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No. US20030091532A1
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Pred. No. 1.9e+06;
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APPLICANT: ANDERSON, KIM V.
APPLICANT: BORNAES, CLAUS
TITLE OF INVENTION: FACTOR VII OR VIIA-LIKE MOLECULES
FILE REFERENCE: 31-001100US
CURRENT APPLICATION NUMBER: US/09/782,587B
CURRENT FILING DATE: 2002-03-26
PRIOR APPLICATION NUMBER: PA 2000 00218
PRIOR FILING DATE: 2000-02-11
PRIOR APPLICATION NUMBER: 60/184,036
PRIOR APPLICATION NUMBER: 60/241,916
PRIOR APPLICATION NUMBER: 60/241,916
PRIOR FILING DATE: 2000-10-18
NUMBER: 05 SEQ ID NOS: 19
COSTENDED: DATE: 2000-10-18
NUMBER: DATE: DATE: 2000-10-18
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FILE REFERENCE: 00-42
CURRENT APPLICATION NUMBER: US/09/892,949
CURRENT FILING DATE: 2001-06-26
PRIOR APPLICATION NUMBER: US 60/214,282
PRIOR APPLICATION NUMBER: US 60/214,955
PRIOR FILING DATE: 2000-06-26
PRIOR PILING DATE: 2000-06-29
PRIOR APPLICATION NUMBER: US 60/267,963
PRIOR APPLICATION NUMBER: US 60/267,963
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US-09-892-949-35
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                                                                                                                                        ; OTHER INFORMATION: FLAG tag peptide sequence US-09-892-949-35
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SEQ ID NO 18
LENGTH: 8
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Matches
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NUMBER OF SEQ ID NOS: 93
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 35
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APPLICANT: Gao, Zeren
APPLICANT: Whitmore, T
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                                                                                                                                                                                ORGANISM: Artificial Sequence FEATURE:
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Kuijper, Joseph L.
Maurer, Mark F.
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Pred. No. 1.9e+06;
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Miltenyi, Stefan
APPLICANT: Miltenyi, Stefan
APPLICANT: Kohler, Matthias
TITLE OF INVENTION: Apparatus and method for modification of
TITLE OF INVENTION: magnetically immobilized biomolecules
FILE REFERENCE: MILT004
CURRENT APPLICATION NUMBER: US/09/994,487
CURRENT FILING DATE: 2001-11-26
NUMBER OF SEQ ID NOS: 7
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2
LENGTH: 8
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OGHER INFORMATION: synthetic peptide
US-09-994-487-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 5, Application US/09803472

Publication No. US20030118990A1

GENERAL INFORMATION:
APPLICANT: Brooks, Alan R.
APPLICANT: Brooks, Alan R.
APPLICANT: Brooks, Alan R.
APPLICANT: Schering Aktiengesellschaft
ITILE OF INVENTION: No. US20030118990A1e1 Estrogen-Regulated G Protein Gamma Subunit:
ITILE OF INVENTION: Compositions and Methods of Use
FILE REFERENCE: 015303-000510US

CURRENT APPLICATION NUMBER: US/09/803,472

CURRENT FILING DATE: 2001-03-10

PRIOR APPLICATION UNUMBER: US 60/188,460

PRIOR APPLICATION NUMBER: US 60/188,460

PRIOR PRIOR APPLICATION VET. 2.1

SEQ ID NO 5

LENGTH: 8
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; ORGANIAM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: General
; OTHER INFORMATION: epitope tag
US-09-803-472-5
Search completed: June 29, 2006, 11:55:57 Job time : 193 secs
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US-09-994-487-2
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Publication No. US20030099954A1
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US-11-313-356-28
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US-11-328-747-16
US-11-328-747-17
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Sequence 13, Application US/09728720A
Publication No. US20060104980A1
GENERAL INFORMATION:
APPLICANT: Foung, Steven K.H.
APPLICANT: Hadlock, Kenneth G.
TITLE OF INVENTION: Antibodies that Inhibit the Interaction of HCV Virions
TITLE OF INVENTION: with Their Receptor.
FILTE OF INVENTION with Their Receptor.
FILE REFERENCE: 2002850-0009
CURRENT APPLICATION NUMBER: US/09/728,720A
CURRENT FILING DATE: 2000-12-01
NUMBER OF SEQ ID NOS: 35
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 13
LENGTH: 8
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Flag Epitope
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US-10-527-411-70
US-10-527-411-71
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US-10-49-902-52167
US-10-49-902-52167
US-10-49-902-52167
US-10-953-349-13783
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                                                  Score 47; DE
Pred. No. 9.6
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24736, Appli

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119209, Appli

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24953, A

24953, A

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35696, A

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35697, A

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35698, A

36988, A
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RESULT 2
US-10-623-808-16
; Sequence 16, Application US/10623808
; Publication No. US20060110802A1
; GENERAL INFORMATION:
; APPLICANT: Alderson, Mark R.
; APPLICANT: Goodwin, Raymond G.
; APPLICANT: Smith, Craig A.
; TITLE OF INVENTION: Cytokine Designated 4-1BB Ligand
; NUMBER OF SEQUENCES: 18
                                                                                                                            RESULT 3
US-10-660-893A-3
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Sequence 3, Application US/10660893A
Publication No. US20060134629A1
GENERAL INFORMATION:
APPLICANT: Link, Charles
TITLE OF INVENTION: Methods and Compositions for Elucidating Protein Expression
TITLE OF INVENTION: Profiles in Cells
FILE REFERENCE: 05237.0003.CPUS00
                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                            IMMEDIATE SOURCE:
CLONE: FLAG peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Anderson, Kathryn A.
REGISTRATION NUMBER: 32,172
REFERENCE/DOCKET NUMBER: 2801-B
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: Apple Power Macintosh
OPERATING SYSTEM: Apple 7.5.3
SOFTWARE: Microsoft Word, Version #6.0.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/623,808
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                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE:
HYPOTHETICAL:
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PRIOR APPLICATION DATA:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/236,918
FILING DATE: 06-May-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
STRANDEDNESS: sir
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE:
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; OTHER INFORMATION: synthesized sequence
US-11-299-013-1
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                                                                                                                                                                                                                                                                                   Sequence 36, Application US/11252276 Publication No. US20060094868A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 1
LENGTH: 8
TYPE: PRT
ORGANISM: Artificial Sequence
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Best Local
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Best Local 9
CURRENT APPLICATION NUMBER: US/11/252,276
CURRENT FILING DATE: 2005-10-17
PRIOR APPLICATION NUMBER: 09/713,572
PRIOR FILING DATE: 2000-11-15
PRIOR APPLICATION NUMBER: 09/430,656
PRIOR FILING DATE: 1999-10-29
PRIOR APPLICATION NUMBER: 09/398,965
                                                                                                                                                                                  APPLICANT: Giuliano, Kenneth A.
APPLICANT: Bright, Gary
APPLICANT: Olson, Keith
APPLICANT: Burroughs-Tencza, S
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APPLICANT: HERNAN, RON
TITLE OF INVENTION: PURIFICATION OF RECOMBINANT PROTEINS FUSED TO MULTIPLE EPITOPES
FILE REFERENCE: SGM6933.7
CURRENT APPLICATION NUMBER: US/11/299,013
CURRENT FILING DATE: 2005-12-09
PRIOR APPLICATION NUMBER: US 09/415,000
PRIOR FILING DATE: 1999-10-08
PRIOR FILING DATE: 1999-10-08
                                                                                                                                        APPLICANT: Burroughs-Tencza, Sarah
TITLE OF INVENTION: A System for Cell Based
FILE REFERENCE: 97-022-K2-CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/660,893A CURRENT FILING DATE: 2003-09-12 NUMBER OF SEQ ID NOS: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US 09/970,308
PRIOR FILING DATE: 2001-10-03
NUMBER OF SEQ ID NOS: 17
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ORGANISM: Artificial
FEATURE:
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Pred. No. 9.6e+04;
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Pred. No. 9.6e+04;
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PRIOR FILING DATE: 1999-09-17

APPLICATION NUMBER: 09/031,271 FILING DATE: 1998-02-27

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                                                                                                                                                                                                                           RESULT 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: Description of Artificial Sequence: FLAG epitope
US-11-252-276-36
             Sequence 5, Application US/11134228A
Publication No. US20060105320A1
GENERAL INFORMATION:
APPLICANT: Lindquist et al.
TITLE OF INVENTION: ELECTRICAL CONDUCTORS AND DEVICES FROM PRION-LIKE PROTEINS
FILE REFERENCE: 30554/40155A
CURRENT APPLICATION NUMBER: US/11/134,228A
CURRENT FILING DATE: 2005-05-20
PRIOR APPLICATION NUMBER: 60/573,277
PRIOR FILING DATE: 2004-05-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 1998-10-30
NUMBER OF SEQ ID NOS: 168
SOFTWARE: PATENTIN VET. 2.0
SEQ ID NO 36
LENGTH: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Wu, Herren
APPLICANT: Go, Changshou
TITLE OF INVENTION: ULTRA HIGH THROUGHPUT CAPTURE LIFT SCREENING METHODS
FILE REFERENCE: AE703US
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 1997-02-27
PRIOR APPLICATION NUMBER: 60/136,078
PRIOR FILING DATE: 1999-05-26
PRIOR APPLICATION NUMBER: 60/106,308
PRIOR APPLICATION NUMBER: 60/106,308
NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
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FILING DATE: 1997-02-27
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8; Conserv
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8; Conservative
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Pred. No.
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; TYPE: PRT
ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: FLAG
US-11-134-228A-5
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Best Local Similarity
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                CURRENT APPLICATION NUMBER: US/11/328,747
CURRENT FILING DATE: 2006-01-09
PRIOR APPLICATION NUMBER: US/10/835,096
PRIOR FILING DATE: 2004-04-28
PRIOR APPLICATION NUMBER: US/09/826,312
PRIOR FILING DATE: 2001-04-03
PRIOR APPLICATION NUMBER: US/09/842,497
PRIOR APPLICATION NUMBER: US/09/542,497
PRIOR APPLICATION NUMBER: US/09/542,497
PRIOR APPLICATION NUMBER: US/09/542,497
PRIOR FILING DATE: 2000-04-03
                                                                                                                                                                                                                                                                                                                                                   Sequence 16, Application US/11328747
Publication No. US20060105409A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn version 3.2 SEQ ID NO 28 LENGTH: 8
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Matches
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PRIOR APPLICATION NUMBER: US 60/286,764
PRIOR FILING DATE: 2001-04-26
NUMBER OF SEQ ID NOS: 80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Lee, Liana
APPLICANT: Krystek, Stanley
APPLICANT: Krystek, Stanley
TITLE OF INVENTION: POLYNUCLEOTIDE ENCODING A NOVEL METALLOPROTEASE HIGHLY EXPRESSED
TITLE OF INVENTION: IN THE TESTIS, MMP-29
FILE REFERENCE: D0141DIV
CURRENT APPLICATION NUMBER: US/11/313,356
CURRENT FILING DATE: 2005-12-21
                                                                                                                                                                                            APPLICANT: Issakani, Sarkiz D.
APPLICANT: Huang, Jianing
APPLICANT: Sheung, Julie
APPLICANT: Sheung, Julie
APPLICANT: Pray, Todd R.
APPLICANT: Rigel Pharmaceuticals, Inc.
APPLICANT: Rigel Pharmaceuticals, Inc.
TITLE OF INVENTION: Ubiquitin Ligase Assay
FILE REFERENCE: 021044-007010US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Wu, Shujian
APPLICANT: Chen, Jian
APPLICANT: Feder, Joh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Ver. 2.0
NUMBER OF
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SEQ ID NOS:
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Feder, John
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RESULT 11

US-11-128-422-16

US-11-128-422-16

Sequence 16, Application US/11128422

Publication No. US20060127985A1

GENERAL INFORMATION:

APPLICANT: Alderson, Mark R.

APPLICANT: Goodwin, Raymond G.

APPLICANT: Smith, Craig A.

TITLE OF INVENTION: Cytokine Designated 4-1BB Ligand NUMBER OF SEQUENCES: 18
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Best Local Similarity
""" Best B; Conserve
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; OTHER INFORMATION: Synthetic epitope
US-11-264-463-2
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SEQ ID NO 2
LENGTH: 8
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CURRENT APPLICATION NUMBER: US/11/264,463
CURRENT FILING DATE: 2005-10-31
PRIOR APPLICATION NUMBER: 60/624,202
PRIOR FILING DATE: 2004-11-01
NUMBER OF SEQ ID NOS: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: BERTOZZI, CAROLYN R.
APPLICANT: AGARD, NICHOLAS J.
APPLICANT: PRESCHER, JENNIFER J.
APPLICANT: BASKIN, JEREMY MICH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: PRESCHER, JENNIFER A.
APPLICANT: BASKIN, JERMY MICHAEL
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
TITLE OF INVENTION: MODIFICATION OF BIOMOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
OTHER INFORMATION: Description of Artificial Sequence:FLAG peptide
OTHER INFORMATION: tag
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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                                                                                                          STREET:
                                        COUNTRY: 1
                                                                                       CITY: Seattle
                                                                                                  ADDRESSEE: Kathryn A. Anderson, Immunex Corporation STREET: 51 University Street
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Pred. No. 9.6e+04;
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Pred. No. 9.6e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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; FEATURE:
; OTHER INFORMATION: synthetic peptide
US-11-270-796-29
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US-11-270-796-29
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Best Local Similarity 1v.
Ches 8; Conservative
                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/11/270,796
CURRENT FILING DATE: 2005-11-08
PRIOR APPLICATION NUMBER: 60/592,592
PRIOR FILING DATE: 2004-07-30
NUMBER OF SEQ ID NOS: 33
SOFTWARE: PatentIn version 3.3
SEQ ID NO 29
                                                     Matches
                                                                 Query Match
Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 29, Application US/11270796
Publication No. US20060134068A1
                                                                                                                                                                                                                                                                                                                    APPLICANT: Dong, Jianli
TITLE OF INVENTION: TREATMENT OF CANCER BY SIMULTANEOUS INHIBITION OF
TITLE OF INVENTION: RESTORATION OR MIMICRY PF p16 Ink4a ACTIVITY
FILE REFERENCE: 02420/1201581-US1
FILE REFERENCE: 02420/1201581-US1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (206) 233-0644
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                           TYPE: PRT
ORGANISM: artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE:
HYPOTHETICAL: N
ANTI-SENSE: NO
                                                                                                                                                                                                      LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IMMEDIATE SOURCE:
CLONE: FLAG peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
PRIOR APPLICATION DATA:
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APPLICATION NUMBER:
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TOPOLOGY: 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: FILING DATE: 07-Ma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
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OPERATING SYSTEM: Apple 7.5.3
SOFTWARE: Microsoft Word, Version #6.0.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
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1 DYKDDDDK 8
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                                                   Similarity
8; Conserv
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                                                   Conservative
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                                               100.0%; Score 47; DB 7; 100.0%; Pred. No. 9.6e+04; tive 0; Mismatches 0;
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100.0%; Pred. No. 9.6e+04;
tive 0; Mismatches 0;
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                Query Match
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                                                                                                   ; OTHER INFORMATION: Synthetic epitope US-11-264-463-3
                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 3, Application US/11264463
Publication No. US20060110782A1
GENERAL INFORMATION:
APPLICANT: BERTOZZI, CAROLYN R.
APPLICANT: AGARD, NICHOLAS J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17, Applica Publication No. US20 GENERAL INFORMATION:
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                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 8
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                            APPLICANT: PRESCHER, JENNIFER A.
APPLICANT: BASKIN, JEREMY MICHAEL
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
TITLE OF INVENTION: MODIFICATION OF BIOMOLECULES
FILE REFERENCE: BERK-039
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US/10/835,096
PRIOR FILING DATE: 2004-04-28
PRIOR APPLICATION NUMBER: US/09/826,312
PRIOR FILING DATE: 2001-04-03
PRIOR APPLICATION NUMBER: US 09/542,497
PRIOR FILING DATE: 2000-04-03
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APPLICANT: Rigel Pharmaceuticals, Inc.
TITLE OF INVENTION: Ubiquitin Ligase Assay
FILE REFERENCE: 021044-007010US
CURRENT APPLICATION NUMBER: US/11/328,747
CURRENT FILING DATE: 2006-01-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Issakani, Sarki
APPLICANT: Huang, Jianing
APPLICANT: Sheung, Julie
                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/11/264,463
CURRENT FILING DATE: 2005-10-31
PRIOR APPLICATION NUMBER: 60/624,202
PRIOR FILING DATE: 2004-11-01
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                                                                                                                                        TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                            LENGTH:
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 1 DYKODDDK 8
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Pray, Todd R.
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                                  Conservative
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Pred. No. 9
                                                  Score 47; DB 7; 1
Pred. No. 9.6e+04;
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                                  Mismatches
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                                                                 SEQ ID NO 26
LENGTH: 20
TYPE: PRT
ORGANISM: Artificial Sequence
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LENGTH: 19
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APPLICANT: MORSE, Clarenc
APPLICANT. YAO, Zhengbin
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SOFTWARE: PatentIn version 3.2
                                                                                                                                          SOFTWARE: PatentIn Ver. 2.1
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ORGANISM: Artificial Sequence
FEATURE:
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Db 1 DYKODDUK 8

RESULT 1:
US-11-251-34-20

US-11-251-34-20

Sequence 20, Application US/1125174

PADDICATION DS. US2006010014A1

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US-11-251-734-18
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Best Local Similarity
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Matches 8; Conservative
                                                        SEQ ID NO 19
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LENGTH: 23
                                                                                                                                       APPLICANT: MORSE, Clarence C
APPLICANT: YAO, Zhengbin
TITLE OF INVENTION: MEMBRANE PENETRATING PEPTIDES AND USES THEREOF
FILE REFERENCE: HMR2053 US NP1
CURRENT APPLICATION NUMBER: US/11/251,734
CURRENT FILING DATE: 2005-10-17
PRIOR APPLICATION NUMBER: US 09/933,780
PRIOR APPLICATION NUMBER: US 09/933,780
PRIOR APPLICATION NUMBER: US 60/227,647
PRIOR APPLICATION NUMBER: US 60/227,647
PRIOR PILING DATE: 2000-08-25
PRIOR FILING DATE: 2000-08-25
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CURRENT APPLICATION NUMBER: US/11/251,734
CURRENT FILING DATE: 2005-10-17
                                                                       NUMBER OF SEQ ID NOS: 54
SOFTWARE: PatentIn versi
                                                                                                            PRIOR APPLICATION NUMBER: GB 0103110.3 PRIOR FILING DATE: 2001-02-07
                                                                                                                                                                                                                                                                                                                                    APPLICANT: AVENTIS PHARMACEUTICALS INC
APPLICANT: GUO, YONG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: GB 0103110.3 PRIOR FILING DATE: 2001-02-07 NUMBER OF SEQ ID NOS: 54
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PRIOR FILING DATE: 2001-08-21
PRIOR APPLICATION NUMBER: US 60/227,647
PRIOR FILING DATE: 2000-08-25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: YAO, Zhengbin
TITLE OF INVENTION: MEMBRANE PENETRATING PEPTIDES AND USES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: AVENTIS PHARMACEUTICALS INC.
APPLICANT: GUO, Yong
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: Synthetic peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence FEATURE:
                                      ENGTH:
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ilarity 100.0%;
Conservative (
                                                                       version 3.2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 47; DB 7;
Pred. No. 0.077;
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APPLICANT: Breung, Julie

APPLICANT: Pray, Todd R.

APPLICANT: Rigel Pharmaceuticals, Inc.

ITILE OF INVENTION: Assays for Identifying Ubiquitin Agents and for TITLE OF INVENTION: Ubiquitin Agents That Modify the Activity of TITLE OF INVENTION: Ubiquitin Agents

FILE REFERENCE: 021044-007061US

CURRENT APPLICATION NUMBER: US/11/090,563

CURRENT FILING DATE: 2005-03-25

PRIOR APPLICATION NUMBER: US 09/542,497

PRIOR APPLICATION NUMBER: US 09/542,497
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Publication No. US20060115864A1
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Best Local
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                                                                                                                                                                                                                                                                                                                     APPLICANT: Issakani, Sarkiz D. APPLICANT: Huang, Jianing
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PRIOR FILING DATE: 2004-04-28
PRIOR APPLICATION NUMBER: US/09/826,312
PRIOR FILING DATE: 2001-04-03
PRIOR PILING DATE: 2000-04-03
PRIOR FILING DATE: 2000-04-03
NUMBER OF SEQ ID NOS: 21
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TYPE: PRT
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Similarity 100.0%;
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Pray, Todd R.
Rigel Pharmaceuticals, Inc.
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Pred. No. 0.3
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Pred. No. 0.077;
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Query Match
Best Local Similarity
Watches 8; Conserva
US-11-090-563-15
; Sequence 15, Application US/11090563
; Publication No. US20060115864A1
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                                                              RESULT 22
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US-11-328-747-15
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US-11-090-563-14
                                                                                                                                                                                                                                                                          US-11-328-747-15
                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 09/542,497
PRIOR FILING DATE: 2000-04-03
NUMBER OF SEQ ID NOS: 21
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 15
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Best Local (
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CURRENT FILING DATE: 2006-01-09
PRIOR APPLICATION NUMBER: US/10/835,096
PRIOR FILING DATE: 2004-04-28
PRIOR FILING DATE: 2004-04-28
PRIOR APPLICATION NUMBER: US/09/826,312
PRIOR FILING DATE: 2001-04-03
PRIOR FILING DATE: 2001-04-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Pray, Todd R.
APPLICANT: Rigel Pharmaceuticals, Inc.
TITLE OF INVENTION: Ubiquitin Ligase Assay
FILE REFERENCE: 021044-007010US
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NUMBER OF SEQ ID NOS: 28
SOFTWARE: PatentIn Ver. 2.1
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PRIOR APPLICATION NUMBER: US 10/108,767
PRIOR FILING DATE: 2002-03-26
PRIOR APPLICATION NUMBER: US 10/109,460
PRIOR FILING DATE: 2002-03-26
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ORGANISM: Artificial Sequence
FEATURE:
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ORGANISM: Artificial Sequence
FEATURE:
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8; Conserv
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                                                                                                                                                                                                             100.0%;
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Pred. No. 0.
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TITLE OF INVENTION: Assays for Identifying Ubiquitin Agents and for TITLE OF INVENTION: Identifying Agents That Modify the Activity of TITLE OF INVENTION: Ubiquitin Agents FILE REFERENCE: 021044-007051US

CURRENT APPLICATION NUMBER: US/11/090,563

CURRENT FILING DATE: 2005-03-25

PRIOR APPLICATION NUMBER: US 09/542,497

PRIOR FILING DATE: 2000-04-03

PRIOR APPLICATION NUMBER: US 09/826,312

PRIOR APPLICATION NUMBER: US 09/826,312

PRIOR APPLICATION NUMBER: US 60/291,863

PRIOR FILING DATE: 2001-05-18

PRIOR FILING DATE: 2001-05-18

PRIOR APPLICATION NUMBER: US 10/091,139

PRIOR APPLICATION NUMBER: US 10/091,174

PRIOR APPLICATION NUMBER: US 10/091,174

PRIOR APPLICATION NUMBER: US 10/091,174

PRIOR APPLICATION NUMBER: US 10/091,174
                                                             APPLICANT: Huang, Jianing
APPLICANT: Sheung, Julie
APPLICANT: Sheung, Julie
APPLICANT: Rigel Pharmaceuticals, Inc.
TITLE OF INVENTION: Ubiquitin Ligase Assay
FILE REFERENCE: 021044-007010US
CURRENT APPLICATION UMBER: US/11/328,747
CURRENT FILING DATE: 2006-01-09
PRIOR APPLICATION NUMBER: US/10/835,096
PRIOR FILING DATE: 2004-04-28
PRIOR FILING DATE: 2004-04-28
PRIOR FILING DATE: 2001-04-03
PRIOR FILING DATE: 2001-04-03
PRIOR APPLICATION NUMBER: US/9542,497
PRIOR FILING DATE: 2000-04-03
PRIOR FILING DATE: 2000-04-03
PRIOR PRIOR DATE: 2000-04-03
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LENGTH: 85
TYPE: PRT
ORGANISM: Artificial Sequence
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SEQ ID NO 18
LENGTH: 86
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PRIOR FILING DATE: 2002-03-26
PRIOR APPLICATION NUMBER: US 10/109,460
PRIOR FILING DATE: 2002-03-26
PRIOR APPLICATION NUMBER: US 10/152,156
PRIOR FILING DATE: 2002-05-20
NUMBER OF SEQ ID NOS: 28
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APPLICANT: Huang, Jianing
APPLICANT: Sheung, Julie
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APPLICANT: Huang, Jianing
APPLICANT: Sheung, Julia
APPLICANT: Pray, Todd R.
APPLICANT: Pray, Todd R.
APPLICANT: Rigel Pharmaceuticals, Inc.
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OTHER INFORMATION: Description of Artificial
OTHER INFORMATION: Sequence:FLAG-Cys-ubiquitin
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APPLICANT: Columbia University
APPLICANT: Angelastro, James M.
APPLICANT: Angelastro, James M.
TITLE OF INVENTION: METHODS FOR REGULATING THE GROWTH AND/OR
TITLE OF INVENTION: EXPRESSION OR FUNCTION OF THE TRANSCRIPTION FACTOR ATFS
FILE REFERENCE: 070050,2879 Greene
CURRENT APPLICATION NUMBER: US/10/971,483
CURRENT APPLICATION NUMBER: US/10/971,483
CURRENT APPLICATION NUMBER: US/089,312
PRIOR APPLICATION NUMBER: US 10/809,312
PRIOR APPLICATION NUMBER: US 60/460,242
PRIOR FILING DATE: 2004-03-24
PRIOR FILING DATE: 2003-04-04
NUMBER OF SEQ ID NOS: 19
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 12
TENCTU: 00
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Publication No. US20060088912A1
GENERAL INFORMATION:
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Best Local S
Matches 8
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                                                                     APPLICANT: CAMPOS, SUSIE J
TITLE OF INVENTION: Compositions and Methods
TITLE OF INVENTION: Glycoprotein (MAG)
FILE REFERENCE: 102729-29
CURRENT APPLICATION NUMBER: US/11/180,855
CURRENT FILING DATE: 2005-07-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: YAN, GOUYING
APPLICANT: XIE, YUHONG
APPLICANT: PAULSEN, JAN
APPLICANT: ZHANG, JIMI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Description of Artificial OTHER INFORMATION: Sequence:FLAG-Ala-Cys-ubiquitin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
APPLICATION NUMBER: US 60/587,893 FILING DATE: 2004-07-14
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PAULSEN, JANE
ZHANG, JIMIN
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BATES, BRIAN
LU, ZHIJIAN
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Conservative Conservati
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100.0%; Pred. No. 0.31;
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Pred. No. 0.36;
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                                                                                                                                                                                                                     of Purifying Myelin-Associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            6; Length 99
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Sequence 6, Application US/11267837A
Publication No. US20060105952A1
GENERAL INFORMATION:
APPLICANT: Allison, Anthony
TITLE OF INVENTION: Modified Annexin Proteins
TITLE OF INVENTION: Transplantation
FILE REFERENCE: SURR.90/CIP2
CURRENT APPLICATION NUMBER: US/11/267,837A
CURRENT FILING DATE: 2005-11-03
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-180-855-2
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US-11-267-837A-6
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PRIOR FILING DATE: 2004-07-15
NUMBER OF SEQ ID NOS: 18
SOFTWARE: PATENTIN VERSION 3.2
SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 2
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Best Local Similarity 100.0%;
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TITLE OF INVENTION: Compositions and Methods of Purifying Myelin-Associated
TITLE OF.INVENTION: Glycoprotein (MAG)
FILE REFERENCE: 102729-29
CURRENT APPLICATION NUMBER: US/11/180,855
CURRENT FILING DATE: 2005-07-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
               PRIOR APPLICATION NUMBER: 60/270,402 PRIOR FILING DATE: 2001-02-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn version 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 18
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PRIOR APPLICATION NUMBER: 11/078,231
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ORGANISM: Homo sapiens
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8; Conserv
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ZHANG, JIMIN
ROOKEY, DIONNA
BATES, BRIAN
LU, ZHIJIAN
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Pred. No. 2
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Pred. No. 1.
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                                                                                                                                                     ; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: FLAG
US-11-134-228A-6
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Publication No. US20060105320A1

GENERAL INFORMATION:

APPLICANT: Lindquist et al.

TITLE OF INVENTION: ELECTRICAL CONDUCTORS AND DEVICES FROM PRION-LIKE PROTEINS

FILE REFERENCE: 30554/40155A

CURRENT APPLICATION NUMBER: US/11/134,228A

CURRENT FILING DATE: 2005-05-20

PRIOR APPLICATION NUMBER: 60/573,277

PRIOR APPLICATION NUMBER: 60/573,277

PRIOR FILING DATE: 2004-05-20

NUMBER: OF SEQ ID NOS: 73
                                                                                                                                                                                                                                 SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 6
LENGTH: 8
TYPE: PRT
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                                                                        Best Local Similarity
Matches 7; Conserv
                                                                                                                Query Match
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PRIOR FILING DATE: 2004-03-11
PRIOR APPLICATION NUMBER: 60/332,582
PRIOR FILING DATE: 2001-11-21
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SOFTWARE: PatentIn version 3.3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc_feature LOCATION: (352)..(352).
LOCATION: (352)..(352) OTHER INFORMATION: The 'Xaa' at location 352 stands for Lys, Asn, Arg, Ser, Thr, OTHER INFORMATION: Ile, Met, Glu, Asp, Gly, Ala, Val, Gln, His, Pro, Leu, Tyr, Trp, OTHER INFORMATION: Cys, or Phe.
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                                                                                                                                                                                                         ORGANISM: Artificial Sequence
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LOCATION: (335)..(335)
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OTHER INFORMATION: T
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                                    1 рукророк в
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8; Conserv
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DYKDEDDK 8
                                                                          Conservative
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, Met, Glu, Asp, Gly, Ala, Val, Gln, His, Pro, Leu, Tyr, Trp,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     'Xaa'
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                                                                                          Score 43; DB 7;
Pred. No. 9.6e+04;
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                                                                          Mismatches
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RESULT 31
US-10-953-349-29832
; Sequence 29832, Application US/10953349
publication No. US20060107345A1
; GENERAL INFORMATION:
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US-11-299-013-17
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US-10-484-105-4
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                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: synthesized inventive sequence US-11-299-013-17
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Publication No. US20060127893A1
GENERAL INFORMATION:
APPLICANT: Universitat Zurich
TITLE OF INVENTION: Modification of Human Variable Domains
FILE REFERENCE: F 2289 US S3
CURRENT APPLICATION NUMBER: US/10/484,105
CURRENT FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: EP 01 11 6756.6
PRIOR FILING DATE: 2001-07-19
NUMBER OF SEQ ID NOS: 19
                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 09/970,308
PRIOR FILING DATE: 2001-10-03
NUMBER OF SEQ ID NOS: 17
SOFTWARE: PatentIn version 3.1
SEQ ID NO 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: BRIZZARD, BILLY L
APPLICANT: HERNAN, RON
TITLE OF INVENTION: PURIFICATION OF RECOMBINANT PROTEINS FUSED TO MULTIPLE EPITOPES
                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17, Application US/11299013 Publication No. US20060088878A1
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CURRENT APPLICATION NUMBER: US/11/299,013
CURRENT FILING DATE: 2005-12-09
PRIOR APPLICATION NUMBER: US 09/415,000
PRIOR FILING DATE: 1999-10-08
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TYPE: PRT
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TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                       FEATURE:
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                                                                                                                           16 DYKDDDD
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100.0%; Prr
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100.0%; Pred. No. 0.7;
Live 0; Mismatches
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Pred. No.
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APPLICANT: ALEXANDROV,

Nickolai et al.

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FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PatentIn version 3.3
SEQ ID NO 29831
LENGTH: 92
                                                             ; TYPE: PRT ; ORGANISM: Zea mays subsp. mays US-10-953-349-33141
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US-10-953-349-33141
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US-10-953-349-29831
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                                                                                                                             Sequence 33141, Application US/10953349

Publication No. US20060107345A1

GENERAL INFORMATION:

APPLICANT: ALEXANDROV, Nickolai et al.

TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES

TITLE OF INVENTION: ENCONDED THERBY

FILE REFERENCE: 2750-1579PUS2

CURRENT FILING DATE: 2004-09-30

NUMBER OF SEQ ID NOS: 40252

SOFTWARE: Patentin version 3.3

SEQ ID NO 33141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: Patentin version 3.3
SEQ ID NO 29932
LEUGTH: 91
TYPE: PRT
ORGANISM: Triticum aestivum
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Best Local Similarity
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Publication NO. US20060107345A1
GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai et al.
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Best Local Similarity
                                   Query Match
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TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
                                                                                                                      LENGTH: 428
                 Local Similarity
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                 Score 39; DE
Pred. No. 29;
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Pred. No. 5.
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Pred. No. 5
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RESULT 36

US-10-953-349-6317

Sequence 6317, Application US/10953349

Publication No. US20060107345A1

GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND COR
TITLE OF INVENTION: ENCONDED THERBY
PILE REFERENCE: 2750-1579FUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
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US-10-953-349-33139

Sequence 33139, Application US/10953349

Publication No. US20060107345A1

GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNJ
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
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US-10-953-349-33140
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US-10-953-349-33139
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Publication No. US20060107345A1

GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai et al.

TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDE:
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTMARE: PARENTIN VERSION 3.3
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Best Local (
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CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
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TYPE: PRT
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Pred. No. 31;
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; ORGANISM: Arabidopsis thaliana
US-10-953-349-6317
RESULT 39
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; ORGANISM: Arabidopsis thaliana
US-10-953-349-6316
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TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: Patentin version 3.3
SEQ ID NO 6316
LENGTH: 294
                                                                                                                                                                                                                  FILE REFERENCE: 2750-15799US2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PatentIn version 3.3
SEQ ID NO 6315
LENGTH: 315
TENTE: DATE: 15799US2
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Publication No. US20060107345A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                   Sequence 6315, Application US/10953349
Publication No. US20060107345A1
GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
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TYPE: PRT
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                                              232 DYNDDDEK 239
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Pred. No.
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Pred. No. 25;
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27;
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RESULT 41
US-10-953-349-21220
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US-10-953-349-21221
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US-10-449-902-39313
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Sequence 21220, Application US/10953349
Publication No. US20060107345A1
GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND
TITLE OF INVENTION: ENCONDED THERBY
TITLE OF INVENTION: ENCONDED THERBY
TITLE OF INVENTION: STOCK THERBY
TITLE OF INVENTION: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
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APPLICANT: ALEXANDROV, Nickolai et al.

TITLE OF INVENTION: SEQUENCE-DEFERMINED DNA FRAGMENTS AND TITLE OF INVENTION: ENCONDED THERBY

FILE REFERENCE: 2750-1579PUSC URRENT APPLICATION NUMBER: US/10/953,349

CURRENT FILING DATE: 2004-09-30
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SEQ ID NO 21221
LENGTH: 389
TYPE: PRT
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
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PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
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TYPE: PRT
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RESULT 43
US-09-756-097B-66
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CURRENT APPLICATION NUMBER: US/09/756,097B
CURRENT FILING DATE: 2001-01-08
PRIOR APPLICATION NUMBER: 09/158,863
PRIOR FILING DATE: 1998-09-23
PRIOR APPLICATION NUMBER: 09/133,717
PRIOR APPLICATION NUMBER: 09/133,717
PRIOR APPLICATION NUMBER: 09/087,233
PRIOR APPLICATION NUMBER: 09/087,233
PRIOR APPLICATION NUMBER: 09/087,233
PRIOR APPLICATION NUMBER: 09/766,354
PRIOR FILING DATE: 1998-08-13
PRIOR FILING DATE: 1996-12-13
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US-10-449-902-45116
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US-10-449-902-45116
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Best Local
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SEQ ID NO 45116
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APPLICANT: Marcia-Blanco, Mariano A.
APPLICANT: Garcia-Blanco, Mariano A.
APPLICANT: Puttaraju, Madaiah
APPLICANT: Mansfield, Gary S.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR USE IN
TITLE OF INVENTION: SPLICEOSOME MEDIATED RNA TRANS-SPLICING IN PLANTS
FILE REFERENCE: A31304-B-A-C 072874,0138
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Publication No. US20060123505A1
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
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Pred. No.
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US-10-449-902-29232
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Publication No. US20060123505A1
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 66
LENGTH: 7
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Best Local (
                  APPLICANT: National Institute of Agrobiological Sciences.
APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
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PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
SOFTWARE: Patentin Ver. 2.1
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
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CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
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TYPE: PRT
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Local Similarity 75.0%;
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Similarity 100.0%; Pred. No.
6; Conservative 0; Mismatcl
SEQ ID NOS:
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GENERAL INFORMATION:

APPLICANT: National Institute of Agrobiological Sciences.

APPLICANT: Bio-oriented Technology Research Advancement Institution.

APPLICANT: Ho Institute of Physical and Chemical Research.

APPLICANT: The Institute of Physical and Chemical Research.

APPLICANT: Foundation for Advancement of International Science.

TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF

FILE REFERENCE: MOA-A0205Y1-US

CURRENT APPLICATION NUMBER: US/10/449,902

CURRENT APPLICATION NUMBER: JP 2002-203269

PRIOR APPLICATION NUMBER: JP 2002-383870

PRIOR FILING DATE: 2002-15-30

PRIOR FILING DATE: 2002-12-11

PRIOR FILING DATE: 2002-12-11
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Best Local Similarity
Watches 6; Conserve
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US-10-449-902-49424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-10-449-902-40889
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US-10-449-902-54492
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                                                                                 CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 40889, Application US/10449902
Publication No. US20060123505A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
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SEQ ID NO 49424
LENGTH: 194
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SEQ ID NO 54492
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                                        SEQ ID NO 40889
LENGTH: 533
                                                                                                                                                                                                                                                                  APPLICANT: National Institute of Agrobiological Sciences.
APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
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ORGANISM: Oryza sativa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            139 DYDDDDDD 145
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85.7%;
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Pred. No. 78;
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APPLICANT: National Institute of Agrobiological Sciences.
APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
FILE REFERENCE: MOA-A020SY1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: UP 2002-203269
PRIOR APPLICATION NUMBER: JP 2002-338870
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
SOFTWARE: PatentIn Ver. 2.1
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CURRENT FILING DATE: 2005-11-03
PRIOR APPLICATION NUMBER: 60/270,402
PRIOR FILING DATE: 2001-02-21
PRIOR APPLICATION NUMBER: 11/078,231
PRIOR FILING DATE: 2005-03-10
PRIOR APPLICATION NUMBER: 10/080,370
PRIOR APPLICATION NUMBER: 10/080,370
PRIOR FILING DATE: 2002-02-21
PRIOR FILING DATE: 2002-02-21
PRIOR APPLICATION NUMBER: 60/552,428
PRIOR APPLICATION NUMBER: 60/579,589
PRIOR APPLICATION NUMBER: 60/579,589
PRIOR FILING DATE: 2004-03-11
PRIOR APPLICATION NUMBER: 60/579,589
PRIOR FILING DATE: 2004-03-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 49
                                                                                                                                                                                                                                                                                                                                                Sequence 34551, Application US/10449902 Publication No. US20060123505A1 GENERAL INFORMATION:
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SOFTWARE: PatentIn versi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Transplantation FILE REFERENCE: SURR.90/CIP2
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TITLE OF INVENTION: Modified Annexin Proteins and Methods for Their Use in Organ
TITLE OF INVENTION: Transplantation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/332,582 PRIOR FILING DATE: 2001-11-21
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ORGANISM: Artificial sequence
FEATURE:
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Pred. No. 9.6e+04;
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Pred. No. 1e+02;
0; Mismatches
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APPLICANT: National Institute of Agrobiological Sciences.

APPLICANT: Bio-oriented Technology Research Advancement Institution.

APPLICANT: The Institute of Physical and Chemical Research.

APPLICANT: Foundation for Advancement of International Science.

ITITLE OF INVENTION: FULL-LENGTH PLANT cDNA AND USES THEREOF FILE REFERENCE: MOA-A0205Y1-US

CURRENT APPLICATION NUMBER: US/10/449,902

CURRENT APPLICATION NUMBER: US/203-05-29

PRIOR APPLICATION NUMBER: JP 2002-203269

PRIOR APPLICATION NUMBER: JP 2002-233870

PRIOR APPLICATION NUMBER: JP 2002-383870

PRIOR APPLICATION NUMBER: JP 3002-383870

PRIOR FILING DATE: 2002-12-11

NUMBER OF SEQ ID NOS: 56791

SOPTWARE: Patentin Ver. 2.1

LENGTH: 342
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; ORGANISM: Oryza sativa
US-10-449-902-40463
                                                                                                                FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
SOFTWARE: Patentin Ver. 2.1
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US-10-449-902-31822
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US-10-449-902-34551
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Matches
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                                                                                              SEQ ID NO 40463
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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Publication No. US20060123505A1
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Best Local Similarity
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Similarity 85.7%;
6; Conservation
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85.7%;
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Pred. No.
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Pred. No.
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                                                                                                                              ; LENGTH: 734
; TYPE: PRT
; ORGANISM: Oryza sativa
US-10-449-902-46241
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US-10-449-902-44996
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                                                                                                                                                                                            PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 46241
LENGTH: 734
                                                           Query Match 74.5%;
Best Local Similarity 85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 44996
LENGTH: 351
                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 46241, Application US/10449902 Publication No. US20060123505A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: National Institute of Agrobiological Sciences.
APPLICANT: Bio-oriented Technology Research Advancement Institute APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
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Publication No. US20060123505A1
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT cDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US-29
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: JP 2002-203269 PRIOR FILING DATE: 2002-05-30
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1 DYKDDDD 7
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Similarity 85.7%;
6; Conservation
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                                         6; Conservative
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85.7%;
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                                         0; Mismatches
                                                             Score 35; DB 6;
Pred. No. 2.1e+02;
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Pred. No.
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                                                                                Length 734;
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RESULT 56
US-10-953-349-6258
; Sequence 6258, Application US/10953349
; Publication No. US20060107345A1
; GENERAL INFORMATION:
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US-10-953-349-39990
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US-10-953-349-18302
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US-10-953-349-39990
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Publication No. US20060107345A1

GENERAL INFORMATION:
APPLICANT: ALEXANDROY, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PatentIn version 3.3
SEQ ID NO 3990
LENGTH: 133
TYCE: DETERMINED THE PARTY OF THE PARTY OF
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TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PatentIn version 3.3
SEQ ID NO 18302
LENGTH: 133
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                            APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
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nes 5; Conserv
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5; Conserv
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PatentIn version 3.3
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Pred. No.
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Pred. No.
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Mismatches
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APPLICANT: National Institute of Agrobiological Sciences.

APPLICANT: Bio-oriented Technology Research Advancement Institution.

APPLICANT: The Institute of Physical and Chemical Research.

APPLICANT: Foundation for Advancement of International Science.

TITLE OF INVENTION: FULL-LENGTH PLANT cDNA AND USES THEREOF

FILE REFERENCE: MOA-A0205Y1-US

CURRENT APPLICATION NUMBER: US/10/449,902

CURRENT TILING DATE: 2003-05-29

PRIOR APPLICATION NUMBER: JP 2002-203269

PRIOR APPLICATION NUMBER: JP 2002-383870

PRIOR FILING DATE: 2002-12-11

NUMBER OF SEQ ID NOS: 56791

SCRIMARDE: Defects user 31
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                                                                   ; ORGANISM: Oryza sativa
US-10-449-902-31740
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US-10-953-349-6257
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Best Local Similarity /1...
Thes 5; Conservative
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APPLICANT: ALEXANDROV, NICKOlai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
                                                                                                                SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 31740
LENGTH: 219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PatentIn version 3.3
SEQ ID NO 6257
LENGTH: 163
                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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Publication No. US20060107345A1
                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 31740, Application US/10449902
Publication No. US20060123505A1
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 Matches
                 Query Match
Best Local
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
                                                                                                     TYPE: PRT
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Similarity 75.06; Conservative
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71.4%;
                 72.3%;
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71.4%;
 Score 34; DB Pred. No. 81; 1; Mismatches
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Pred. No.
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RESULT 61
US-10-953-349-2040
US-10-953-349-2040
; Sequence 2040, Application US/10953349
; Publication No. US20060107345A1
; GENERAL INFORMATION:
; APPLICANT: ALEXANDROV, Nickolai et al.
; APPLICANT: ALEXANDROV, Nickolai et al.
; TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
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US-10-953-349-6256
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US-10-449-902-40441
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CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40552
SOFTWARE: PatentIn version 3.3
SEQ ID NO 6256
LENGTH: 222
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Publication No. US20060123505A1
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TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: JS 2003-05-29
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 2750-1579PUS2
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PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 40441
LENGTH: 219
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75.0%;
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Pred. No. 81;
1; Mismatches
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PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
PRIOR FPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 41697
ENOTH: 294
US-10-953-349-36044
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US-10-953-349-36044
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US-10-449-902-41697
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Publication No. US20060107345A1
GENERAL INFORMATION:
                                                                                    SEQ ID NO 36044
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CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PatentIn version 3.3
SEQ ID NO 2040
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Best Local :
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                                                                                                                                           APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES.
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
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                                                                                                     NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PatentIn version
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL LENGTH PLANT CDNA AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
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             ORGANISM: Zea mays subsp. mays
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TYPE: PRT
                                                           LENGTH: 301
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71.4%;
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75.0%;
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; Pred. No. 85;
2; Mismatches
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Pred. No. 1.1e+02;
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CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PatentIn version 3.3
SEQ ID NO 2039
LENGTH: 338
TYPE: PRT
ORGANISM: Arabidopsis thaliana
US-10-953-349-2039
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Best Local Similarity
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Best Local Similarity
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Best Local Similarity
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US-10-953-349-5131
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US-10-953-349-2039
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TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT PELLORATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: Patentin version 3.3
SEQ ID NO 2038
LENGTH: 340
TYPE: PRT
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GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750.01579PUS2
CITETREE REFERENCE: 2750.01579PUS2
                             Sequence 5131, Application US/10953349 Publication No. US20060107345A1 GENERAL INFORMATION:
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TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
                APPLICANT: ALEXANDROV, Nickolai et al.
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85.7%;
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Pred. No. 1.3e+02;
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Pred. No. 1.3e+02;
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Pred. No. 1.1e+02;
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US-10-449-902-52372
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APPLICANT: National Institute of Agrobiological Sciences.
APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
APPLICANT: Foundation for Advancement of International Science.
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CURRENT FILING NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: Patentin version 3.3
SEQ ID NO 5131
LENGTH: 362
TYPE: PRT
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Publication No. US20060123505A1
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LENGTH: 373
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Best Local (
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APPLICANT: Bio-oriented Technology Research Advancement Institute APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
CURRENT FILING DATE: 2003-05-29
                 FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
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PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
SOFTWARE: Patentin Ver. 2.1
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TYPE: PRT
NUMBER OF SEQ ID NOS: 56791
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71.4%;
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Pred. No. 1.4e
2; Mismatches
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Pred. No. 1.4e+02;
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Sequence 42067, Application US/10449902
Publication No. US20060123505A1
GENERAL INFORMATION:
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US-10-449-902-42067
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CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
SOFTWARE: Patentin Ver. 2.1
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; ORGANISM: Oryza sativa
US-10-449-902-43294
                                                                 SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 50266
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Publication No. US20060123505A1
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Best Local
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT cDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER: OF SEQ ID NOS: 56791
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APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL LENGTH PLANT CDNA AND USES THEREOF
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APPLICANT: Bio-oriented Technology Research Advancement
APPLICANT: The Institute of Physical and Chemical Resear
APPLICANT: Foundation for Advancement of International S
ORGANISM: Oryza sativa
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Pred. No. 1.7e+02
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Pred. No. 1.6e+02;
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; ORGANISM: Oryza sativa
US-10-449-902-45621
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US-10-449-902-44569
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Query Match
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SEQ ID NO 45621
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Publication No. US20060123505A1
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
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                                                                                                                                                            LENGTH: 521
TYPE: PRT
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TYPE: PRT
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71.4%;
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Pred. No. 2e+02;
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Pred. No. 2e+02;
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Pred. No. 1.9e+02
2; Mismatches
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US-10-953-349-21280
US-10-953-349-21280, Application US/10953349
Sequence 21280, Application US/10953349
; Publication No. US20060107345A1
; GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCOUNDED THERBY
FILE REFERENCE: 2750-1579PUS2
; CURRENT APPLICATION NUMBER: US/10/953,349
; CURRENT FILING DATE: 2004-09-30
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                   US-10-449-902-30504
; Sequence 30504, Application US/10449902
; Publication No. US20060123505A1
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US-10-449-902-45113
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US-10-449-902-45113
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APPLICANT: National Institute of Agrobiological Sciences.
APPLICANT: Bio-oriented Technology Research Advancement
APPLICANT: The Institute of Physical and Chemical Reseat
APPLICANT: Foundation for Advancement of International S
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SEQ ID NO 45113
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GENERAL INFORMATION:
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Matches 6; Conserv
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APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
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PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
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SOFTWARE: PatentIn version 3.3
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nes 5; Conserv
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71.4%;
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75.0%;
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Pred. No. 2.7e+02;
Pred. No. 2.7e+02;
Indels
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Pred. No. 37;
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT cDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
SOFTWARE: Patentin Ver. 2.1
SOFTWARE: Patentin Ver. 2.1
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US-10-449-902-30504
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US-10-449-902-34122
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                     GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai et al.
APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
                                                                                                                                                  Sequence 27604, Application US/10953349
Publication No. US20060107345A1
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Best Local Similarity
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APPLICANT: Bio-oriented Technology Research Advancement
APPLICANT: The Institute of Physical and Chemical Resear
APPLICANT: Foundation for Advancement of International S
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APPLICANT: The Institute of Physical and Chemical Research
APPLICANT: Foundation for Advancement of International Science.
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CURRENT APPLICATION NUMBER: US/10/953,349
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TYPE: PRT
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85.7%;
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Pred. No.
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APPLICANT: National Institute of Agrobiological Sciences.

APPLICANT: Bio-oxiented Technology Research Advancement Institution.

APPLICANT: The Institute of Physical and Chemical Research.

APPLICANT: Foundation for Advancement of International Science.

ITITLE OF INVENTION: FULL-LENGTH PLANT cDNA AND USES THEREOF FILE REFERENCE: MOA-0205Y1-US

CURRENT APPLICATION NUMBER: US/10/449,902

CURRENT FILING DATE: 2003-05-29

PRIOR APPLICATION NUMBER: JP 2002-203269

PRIOR APPLICATION NUMBER: JP 2002-383870

PRIOR APPLICATION SUMBER: JP 2002-383870

PRIOR PRIVING DATE: 2002-12-11

NUMBER OF SEG ID NOS: 56791

SOFTWARE: PatentIn Ver. 2.1
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US-10-449-902-30872
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US-10-449-902-50940
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                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 30872, Application US/10449902
Publication No. US20060123505A1
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SEQ ID NO 50940
LENGTH: 221
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Best Local :
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NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PatentIn version 3.3
SEQ ID NO 27604
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
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TYPE: PRT
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les 6; Conserv
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Local Similarity 85.7%;
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85.7%;
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Pred. No. 1.2e+02;
0; Mismatches 1
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Pred. No. 1.1e+02;
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APPLICANT: CHIRON SPA
TITLE OF INVENTION: STAPHYLOCOCCUS AUREUS PRO
FILE REFERENCE: P026927WO
CURRENT APPLICATION NUMBER: US/10/471,571A
CURRENT FILING DATE: 2003-09-12
PRIOR APPLICATION NUMBER: GB-0107661.1
PRIOR FILING DATE: 2001-03-27
NUMBER OF SEQ ID NOS: 5642
SOFTWARE: SegWin99, version 1.03
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US-10-471-571A-3230
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Best Local Similarity
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US-10-449-902-30872
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APPLICANT: National Institute of Agrobiological Sciences.

APPLICANT: Bio-oriented Technology Research Advancement Institution.

APPLICANT: The Institute of Physical and Chemical Research.

APPLICANT: Foundation for Advancement of International Science.

TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF

FILE REFERENCE: MOA-A0205Y1-US

CURRENT APPLICATION NUMBER: US/10/449,902

CURRENT FILING DATE: 2003-2007-20026
                                                                                                                                                                                                            SEQ ID NO 3230
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SEQ ID NO 37911
                    Query Match
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Publication No. US20060123505A1
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
                                                                                                                              LENGTH: 553
TYPE: PRT
ORGANISM: Staphylococcus aureus
FEATURE:
                                                                    NAME/KEY: MISC_FEATURE
LOCATION: (1)...(553)
OTHER INFORMATION: arginyl-trna synthetase
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71.4%;
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75.0%;
70.2%;
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Score 33; DB 6;
Pred. No. 3.1e+02;
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Pred. No.
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Pred. No. 2.
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2.8e+02;
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                Length 553;
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Best Local Similarity
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US-10-449-902-31438
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US-10-449-902-40941
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 31438
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Publication No. US20060123505A1
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Best Local Similarity 62.5%;
Matches 5; Conservative
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LENGTH: 770
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CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
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PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
SOFTWARE: Patentin Ver. 2.1
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818 YKDDEDE 824
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Pred. No. 4.4e+02;
                                                                                                                                                                              Score 33; DB 6;
Pred. No. 5e+02;
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US-10-449-902-52762
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Best Local Similarity
Thes 5; Conserva
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CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: UP 2002-203269
PRIOR APPLICATION NUMBER: UP 2002-203269
PRIOR FILING DATE: 2002-05-30
PRIOR FILING DATE: 2002-05-30
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 56037
                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 52762
LENGTH: 1179
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Publication No. US20060123505A1
                                                                                           Query Match 70.3
Best Local Similarity 62.9
Matches 5; Conservative
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT cDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US
CURRENT APPLICATION NUMBER: JP 2002-203269
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
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APPLICANT: Bio-oriented Technology Research Advancement
APPLICANT: The Institute of Physical and Chemical Resear
APPLICANT: Foundation for Advancement of International S
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PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
                                                                                                                                                                                                                                              TYPE: PRT
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87 EYDDDDDDE 94
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71.4%;
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Pred. No. 5.4e+02;
                                                                                                                       Score 33; DB 6;
Pred. No. 6.9e+02;
                                                                                                  Mismatches
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RESULT 86 US-10-449-902-53947

; Sequence 53947, Application US/10449902

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US-11-289-102-378
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US-10-449-902-53947
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CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
                                                                                                        GENERAL INFORMATION:
                                                                                                                          Sequence 41, Application US/11074497 Publication No. US20060122135A1
                                                                                                                                                                                                                                                                                                                                     Matches
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SEQ ID NO 378
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SEQ ID NO 53947
LENGTH: 1180
               APPLICANT: Geerts, Hugo
APPLICANT: Masure, Stefan
APPLICANT: Cik, Miroslav
APPLICANT: Meert, Theo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: BIOMARKERS AND METHODS FOR DETERMINING SENSITIVITY TITLE OF INVENTION: MICROTUBULE-STABILIZING AGENTS FILE REPERENCE: 10338 NP CURRENT APPLICATION NUMBER: US/11/289,102 CURRENT FILING DATE: 2005-11-29 PRIOR APPLICATION NUMBER: US 60/631,993 PRIOR APPLICATION NUMBER: US 60/631,993 PRIOR FILING DATE: 2004-11-30 NUMBER OF SEQ ID NOS: 395
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Lee, Hyerim APPLICANT: Shaw, Peter APPLICANT: Clark, Edw:
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
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les 5; Conserv
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Clark, Edwin
Ver Donck, Luc
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71.4%;
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71.4%;
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Pred. No. 7.2e+02;
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Pred. No. 6.9e+02;
2; Mismatches 0
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RESULT 90
US-10-449-902-33753
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US-10-449-902-43449
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CURRENT FILING DATE: 2005-03-08
PRIOR APPLICATION NUMBER: US/09/357,349
PRIOR FILING DATE: 1999-07-14
PRIOR APPLICATION NUMBER: 09/327,668
PRIOR FILING DATE: 1999-06-08
PRIOR APPLICATION NUMBER: 09/248,772
PRIOR APPLICATION NUMBER: 09/248,772
PRIOR FILING DATE: 1999-02-12
PRIOR APPLICATION NUMBER: GB 9815283.8
PRIOR FILING DATE: 1999-07-14
NUMBER OF SEQ ID NOS: 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: National Institute of Agrobiological Sciences.
APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
FILE REFERENCE: NOA-A0205Y1-US
CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: US/10/449,902
PRIOR APPLICATION NUMBER: US/10/203269
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR RILING DATE: 2002-05-30
PRIOR FILING DATE: 2002-05-30
PRIOR FILING DATE: TO THE TECHNOLOGY TO THE TECHN
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SEQ ID NO 43449
LENGTH: 74
Sequence 33753, Application US/10449902
Publication No. US20060123505A1
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Best Local (
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Publication No. US20060123505A1
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LENGTH: 39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
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                                                                                                                                                                                                                                            55 YKDEDD 60
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5; Conservative
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75.0%;
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Pred. No. 52;
1; Mismatches
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Pred. No. 26;
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GENERAL INFORMATION

APPLICANT: National Institute of Agrobiological Sciences.
APPLICANT: Bio-oriented Technology Research Advancement Inc
APPLICANT: The Institute of Physical and Chemical Research

Institution.

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Query Match
Best Local Similarity
Matches 5; Conserva
                                                                                                                                                                    RESULT 92
US-10-953-349-6227
; Sequence 6227, Application US/10953349
; Publication No. US20060107345A1
; GENERAL INFORMATION:
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US-10-449-902-36276
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CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
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SEQ ID NO 36276
LENGTH: 146
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
                 APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: ENCONDED THERBY
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION INUMBER: US/10/953,349
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
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CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
PRIOR FILING DATE: 2002-05-30
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                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity nes 5; Conserv
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PatentIn version 3.3
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83.3%;
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71.4%;
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Pred. No. 1.1e+02;
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Pred. No. 56;
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Query Match
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                                                                                                              FILE REFERENCE: 270.78USWO
CURRENT APPLICATION NUMBER: US/10/516,753
CURRENT FILING DATE: 2004-12-03
PRIOR APPLICATION NUMBER: PCT/CA2003/000822
PRIOR FILING DATE: 2003-06-03
PRIOR APPLICATION NUMBER: US 60/387,088
PRIOR FILING DATE: 2002-06-06
NUMBER: OF SEQ ID NOS: 104
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US-10-449-902-29282
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US-10-449-902-29282
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LENGTH: 203
                                                       SOFTWARE: PatentIn version 3.0 SEQ ID NO 74 LENGTH: 221
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SEQ ID NO 29282
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: National Institute of Agrobiological Sciences.
APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
                                                                                                                                                                                                                                                                                     APPLICANT: Hannoufa, Abdelali
APPLICANT: Lydiate, Derek J.
APPLICANT: Gao, Ming-Jun
TITLE OF INVENTION: REGULATION OF GENE EXPRESSION USING CHROMATIN REMODELLING FACTOR'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 205
TYPE: PRT
TYPE: PRT
ORGANISM: Arabidopsis
FEATURE:
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75.0%;
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Pred. No. 1.5e+02;
0; Mismatches 1
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Pred. No. 1.5e+02
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OTHER INFORMATION: atKCL2

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US-10-953-349-16218; Sequence 16218, Application US/10953349; Publication No. US20060107345A1; GENERAL INFORMATION: ALEXANDROV, Nickolai et al.
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                                                                                        RESULT 97
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                                                                                                                                                                                                                                                                               ; ORGANISM: Glycine max US-10-953-349-16219
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US-10-953-349-6226
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Best Local S
Matches 5
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TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES

TITLE OF INVENTION: ENCONDED THERBY

FILE REFERENCE: 2750-1579PUS2

CURRENT APPLICATION NUMBER: US/10/953,349

CURRENT FILING DATE: 2004-09-30

NUMBER OF SEQ ID NOS: 40252

SOFTWARE: PatentIn version 3.3

SEQ ID NO 16219

LENGTH: 234
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Publication No. US20060107345A1
GENERAL INFORMATION:
APPLICANT: ALEXANDROY, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOOTWARE: Patentin version 3.3
SEQ ID NO 6226
LENGTH: 231
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Publication No. US20060107345A1
GENERAL INFORMATION:
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Local Similarity 83.3%;
les 5; Conservative
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71.4%;
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75.0%;
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Pred. No. 1.7e+02
0; Mismatches
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Pred. No. 1.8e+02;
1; Mismatches 1;
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Pred. No.
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1.7e+02;
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                                                                                 ; ORGANISM: Glycine max US-10-953-349-20549
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US-10-953-349-31660
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                                                                                                                 Sequence 20549, Application US/10953349

Publication No. US20060107345A1

GENERAL INFORMATION:
APPLICANT: ALEXANDROY, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES,
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349

CURRENT FILING DATE: 2004-09-30

NUMBER OF SEQ ID NOS: 40252
SOFTWARE: Patentin version 3.3

SEQ ID NO 20549
LENGTH: 255
TYPE: PRT
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Publication No. US20060107345A1

GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDE:
TITLE OF INVENTION: ENCONDED THERBY
FILLE REFERENCE: 2750-1579FUS2
CURRENT APPLICATION UNMERR: US/10/953,349

CURRENT FILING DATE: 2004-09-30

NUMBER OF SEQ ID NOS: 40252
SOFTWARE: Patentin version 3.3

SEQ ID NO 31660

LENGTH: 252
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TITLE OF INVENTION: ENCONDED THERBY
FILE REPERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOPTWARE: Patentin version 3.3
SEQ ID NO 16218
LENGTH: 240
TYPE: PRT
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Best Local Similarity 71.4%;
Matches 5; Conservative
Query Match
Best Local Similarity 71.4
Matches 5; Conservative
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Best Local :
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Similarity 62.5%;
5; Conservative
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                     68.1%;
71.4%;
Score 32; DB 6; Length 255; Pred. No. 1.9e+02; 1; Mismatches 1; Indels
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Pred. No. 1.9e+02;
1; Mismatches 2;
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Pred. No. 1.8e+02;
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RESULT 100
US-10-953-349-31659
Sequence 31659, Application US/10953349
Publication No. US20060107345A1
GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579FUS2
CURRENT APPLICATION UNMERR: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PATENTIN VERSION 3.3
SEQ ID NO 31659
LENGTH: 260
TYPE: PRT
ORGANISM: Triticum aestivum
US-10-953-349-31659
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Search completed: June 29, 2006, 11:56:19 Job time : 25 secs
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Best Local Similarity 62.5%;
Matches 5; Conservative
                                                                        1 DYKDDDDK 8
|| |||:
147 DYSSDDDE 154
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Pred. No. 2e+02;
1; Mismatches
                                                                                                                                                 2; Indels
                                                                                                                                                                                  Length 260;
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